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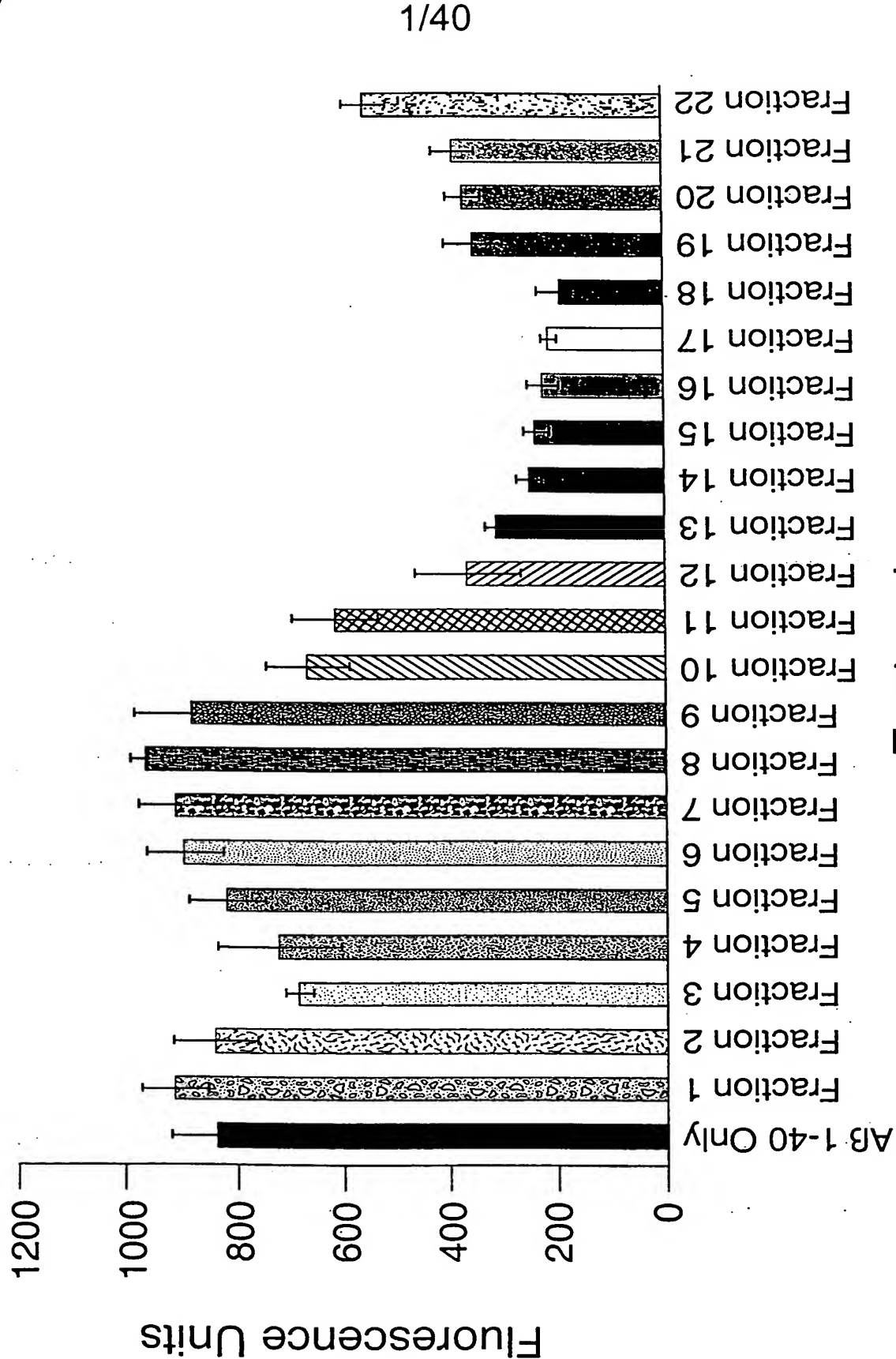


Fig. 1

Treatment

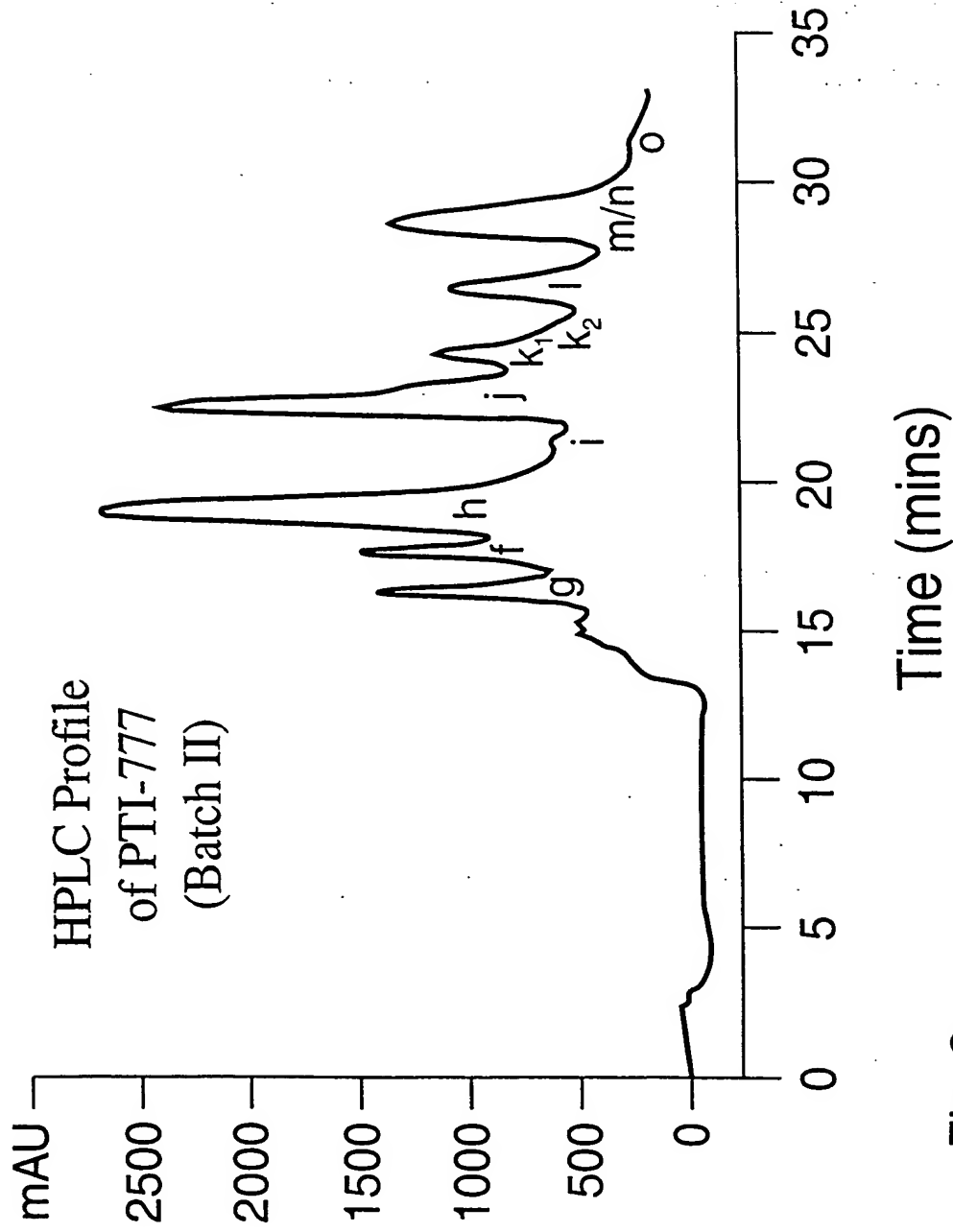


Fig. 2

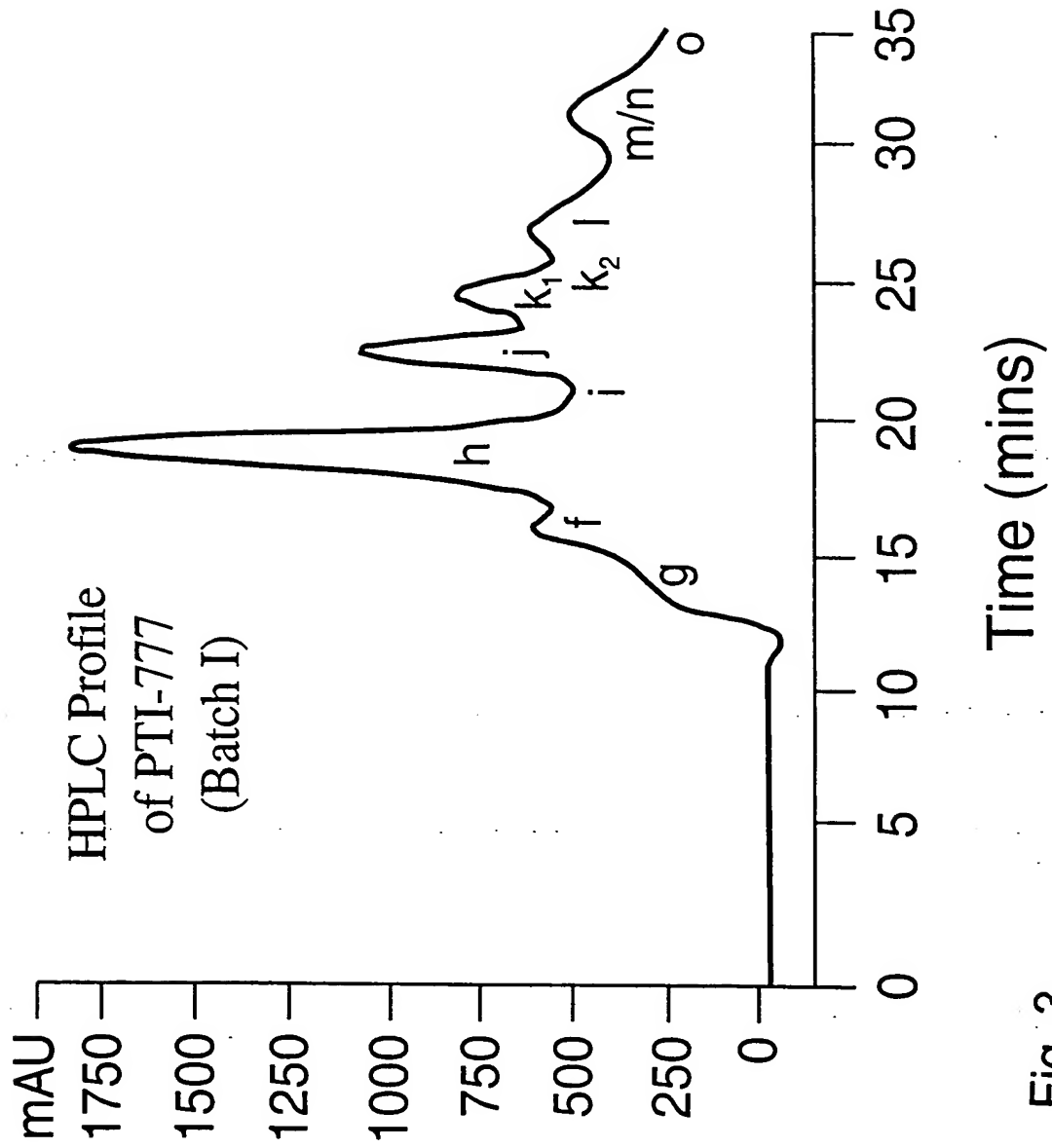
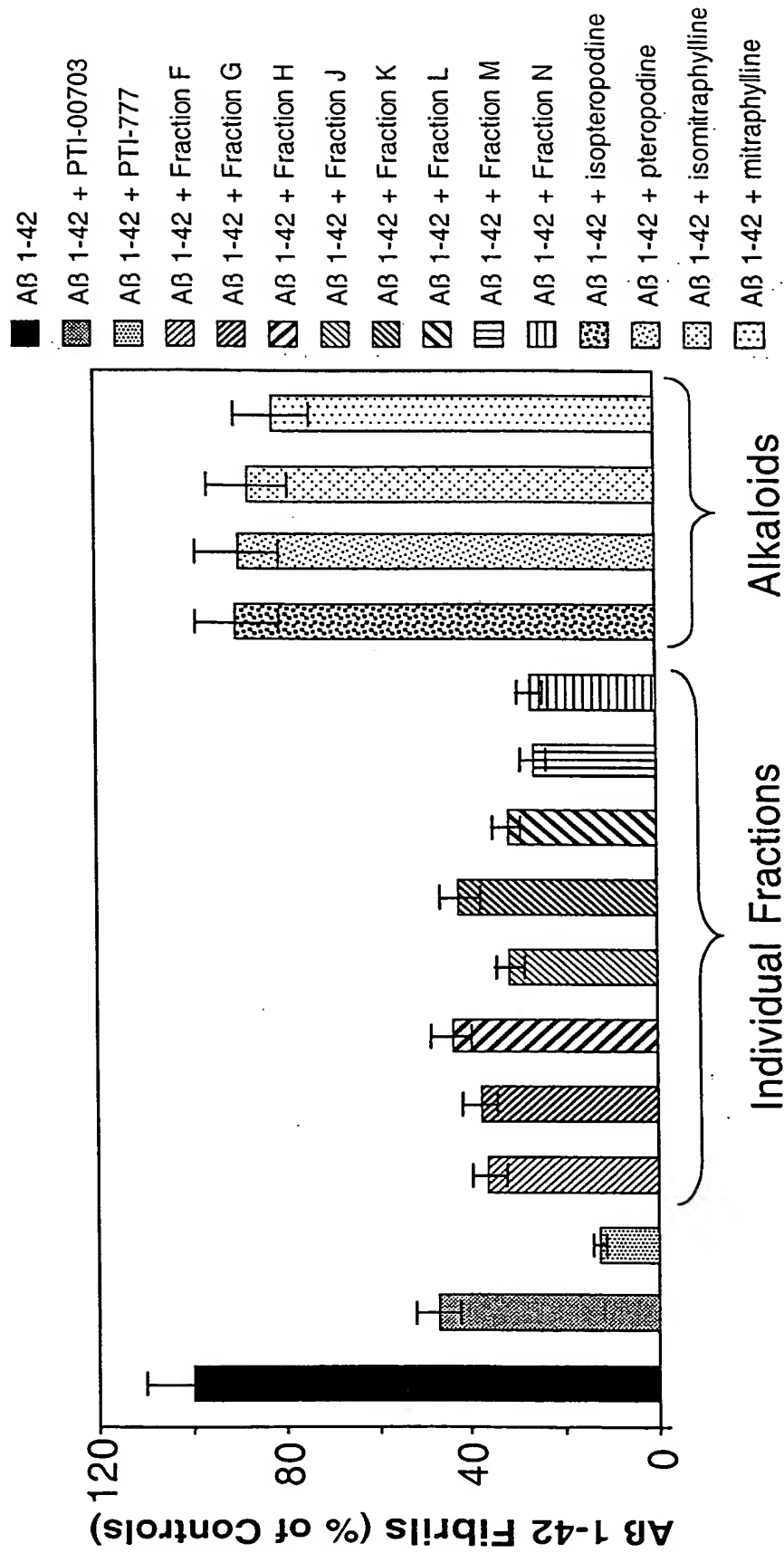


Fig. 3



Treatment Groups

Fig. 4

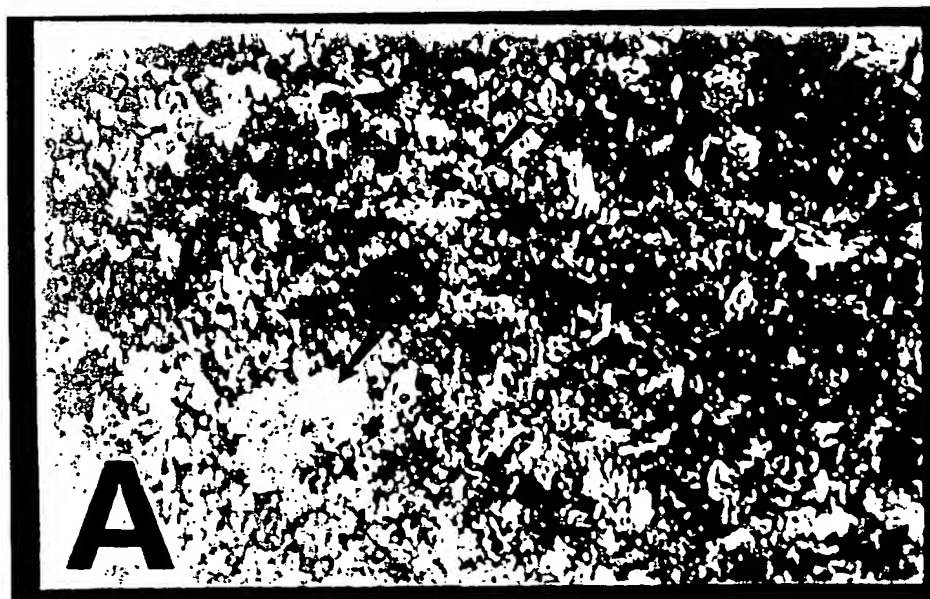


Fig. 5A

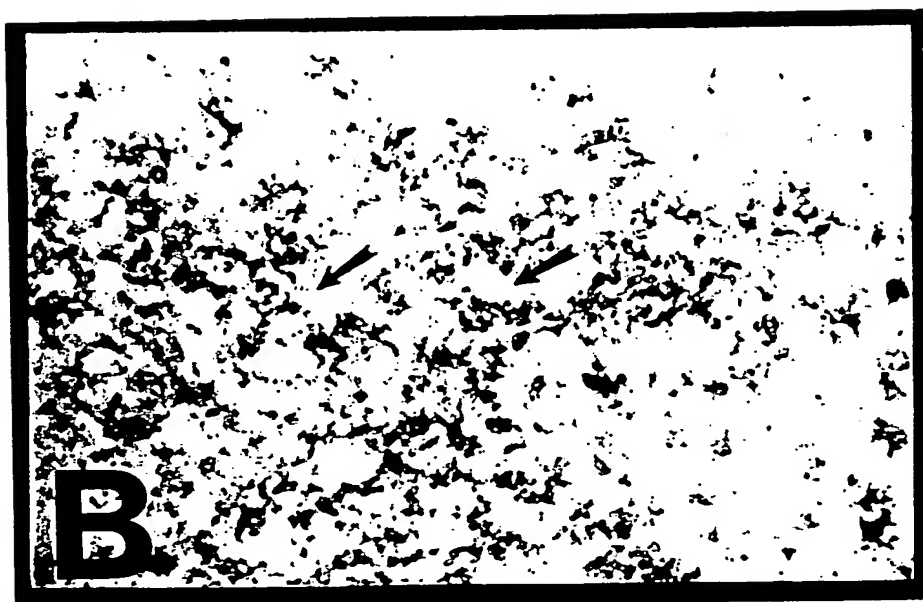


Fig. 5B



Fig. 5C



Fig. 5D

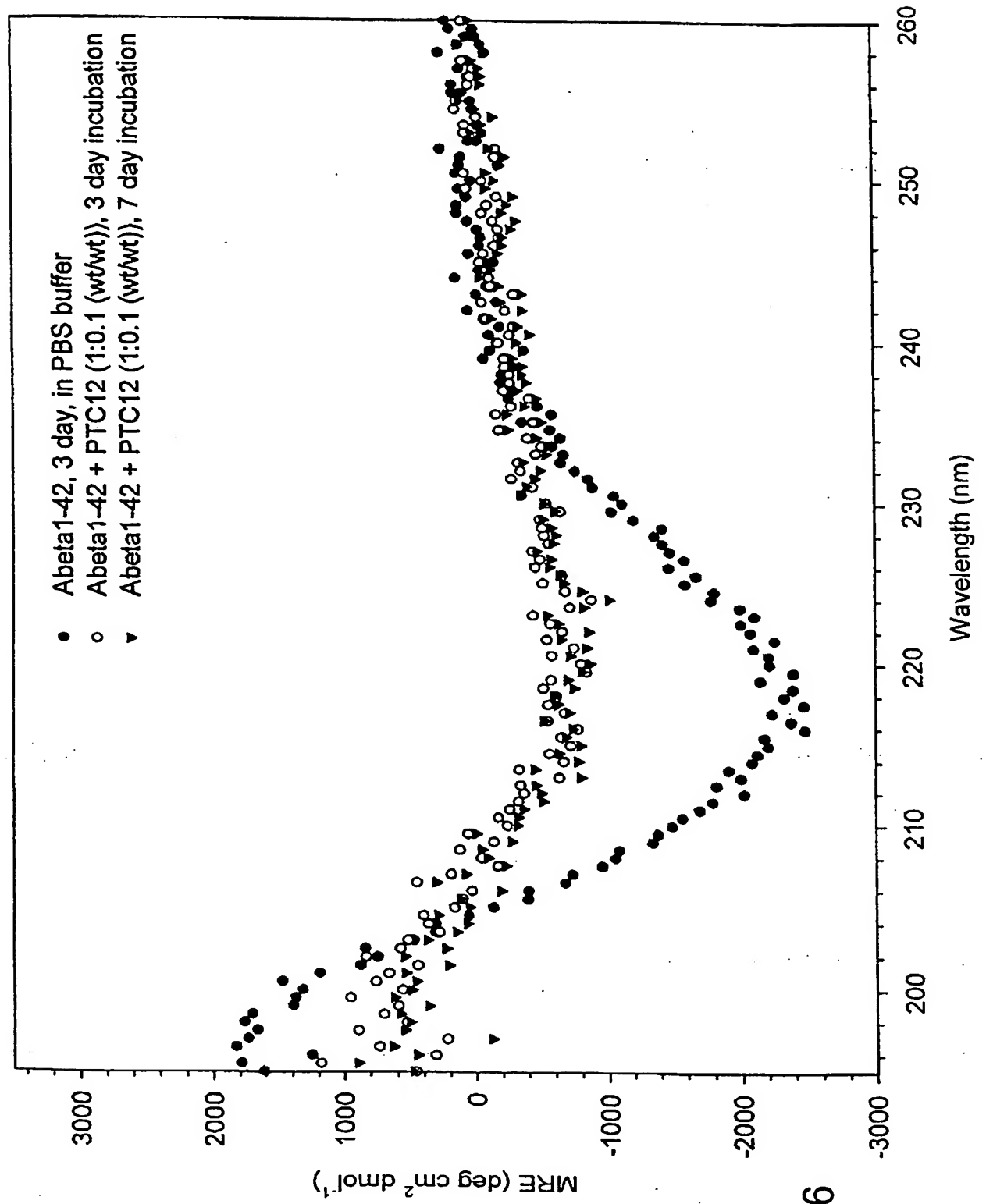


Fig. 6

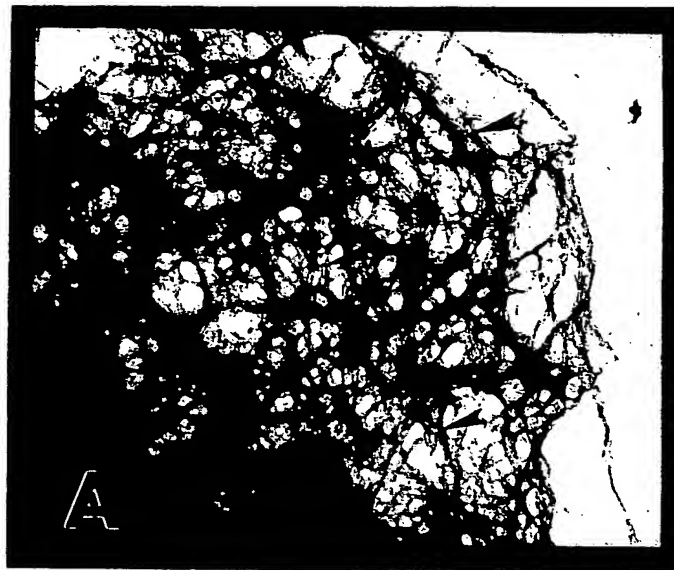


Fig. 7A

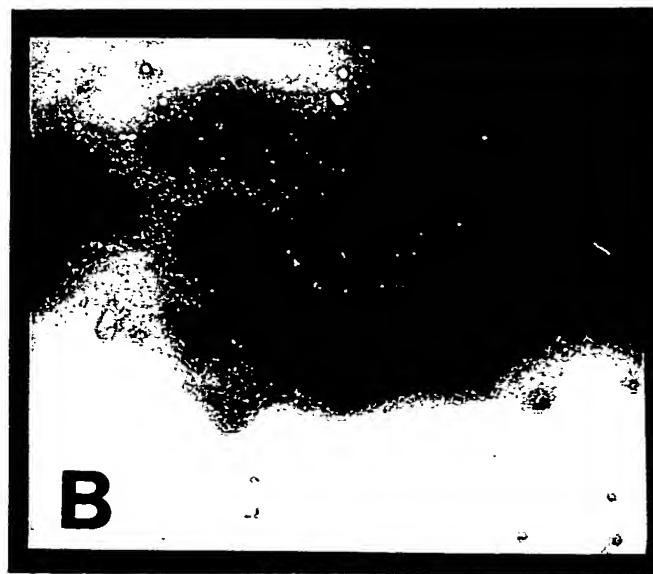


Fig. 7B

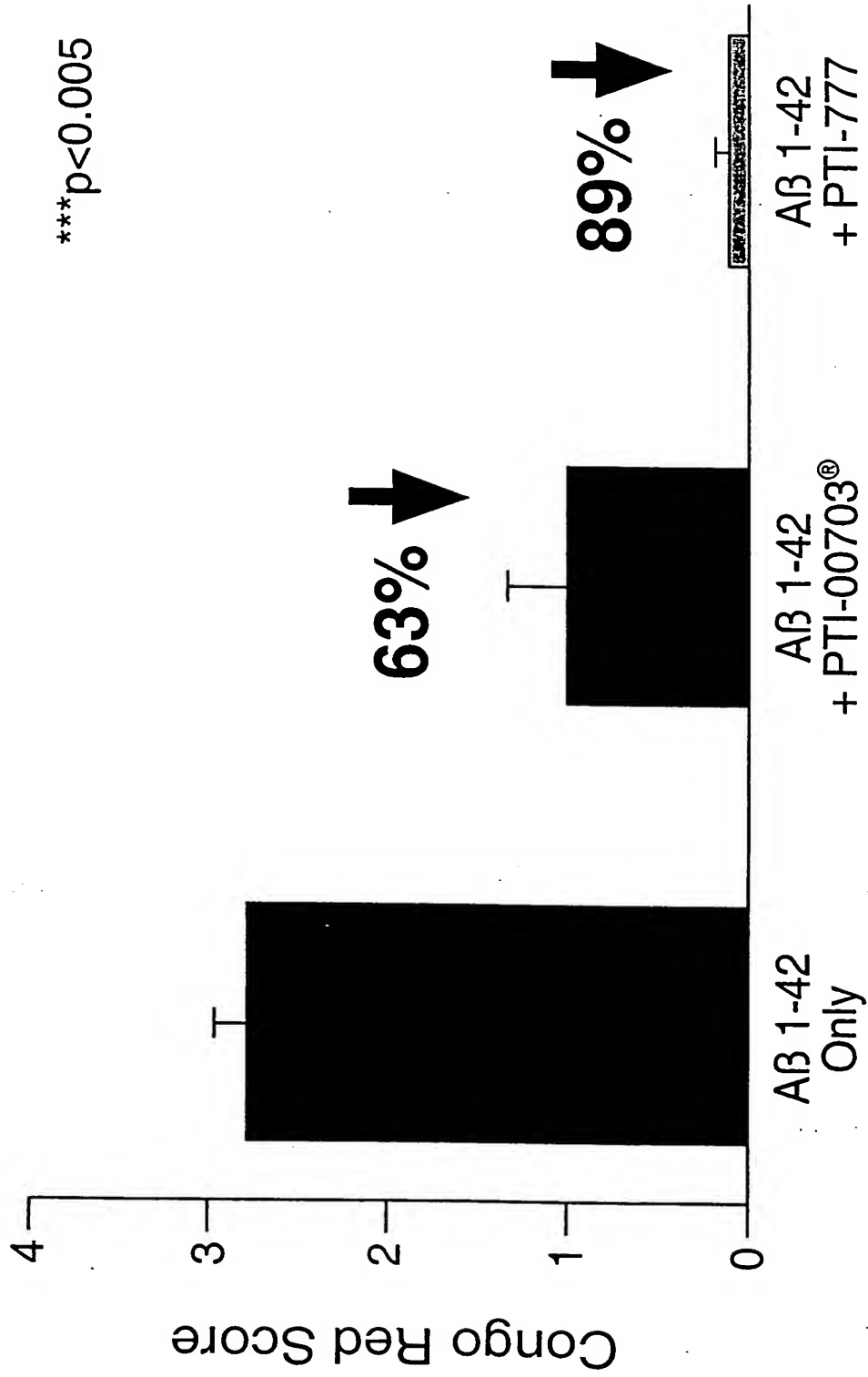


Fig. 8

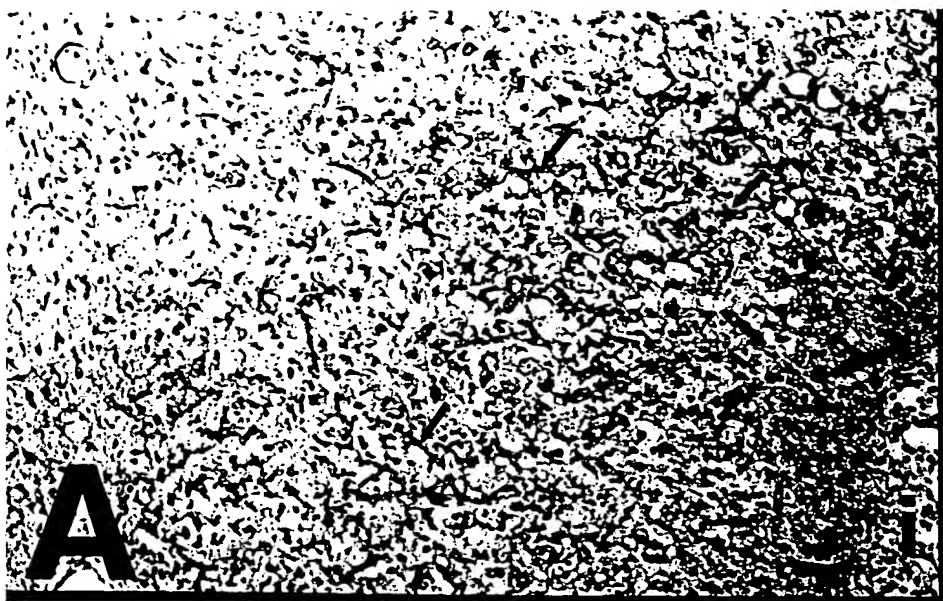


Fig. 9A

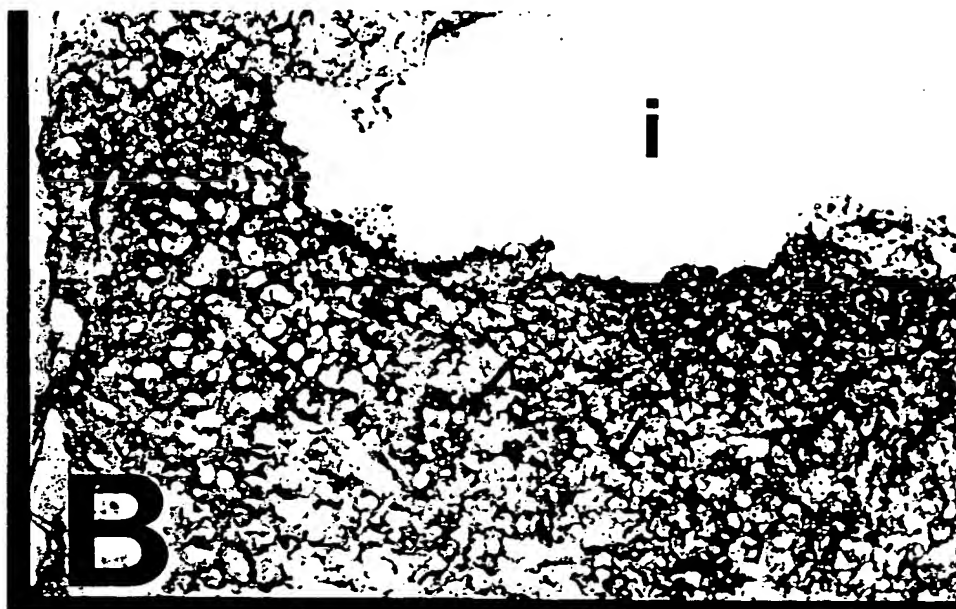


Fig. 9B

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Fig. 9C



Fig. 9D

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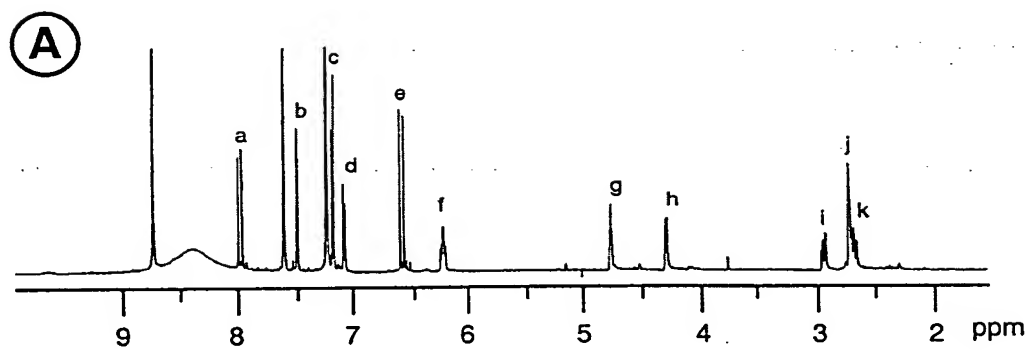


Fig. 10A

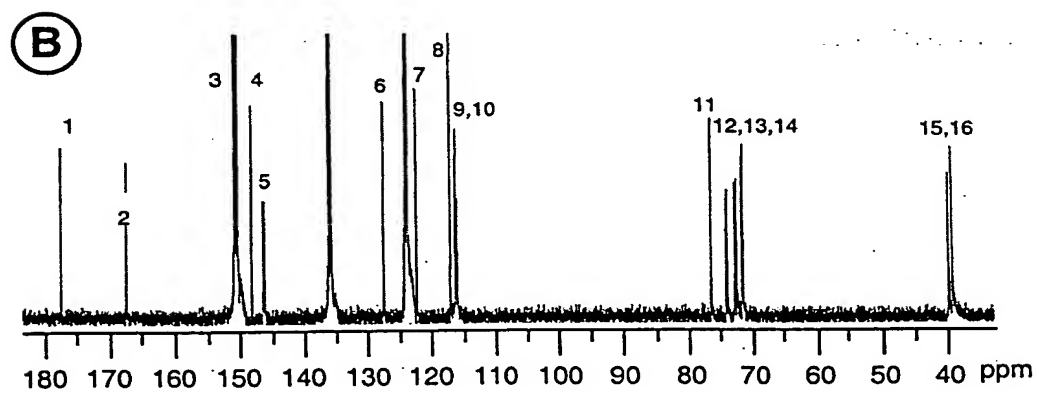


Fig.10B

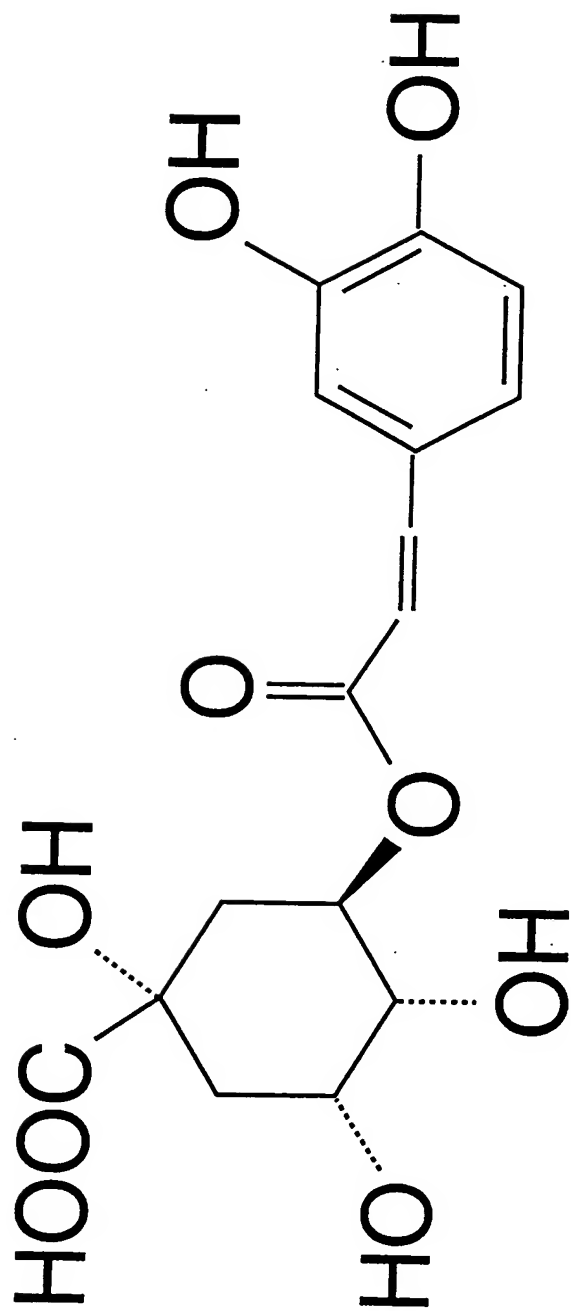


Fig. 11

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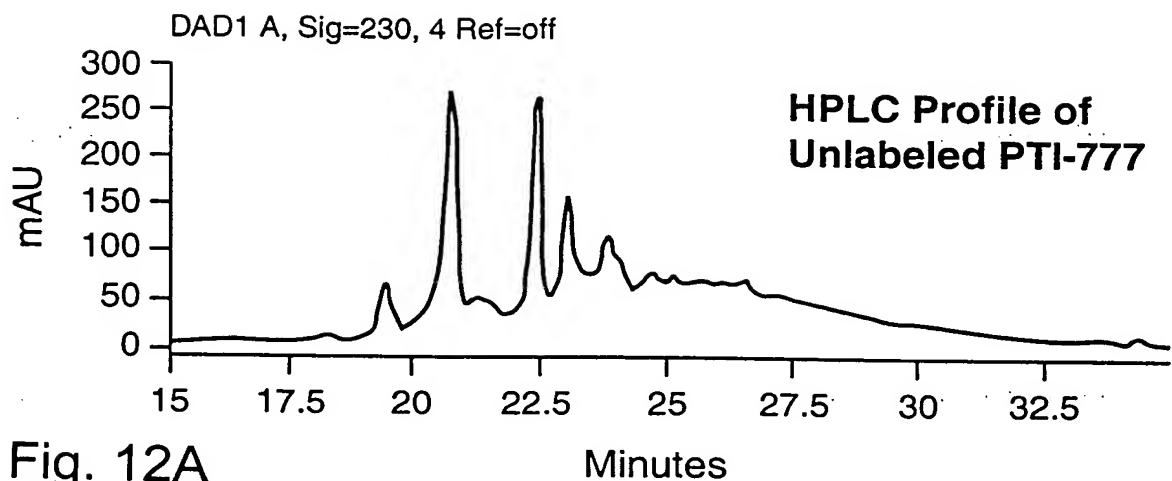


Fig. 12A

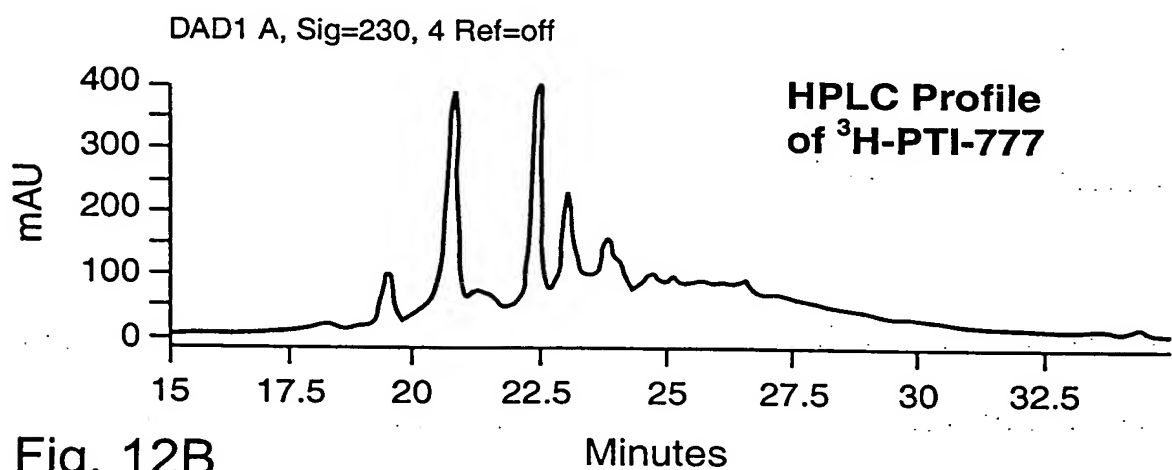


Fig. 12B

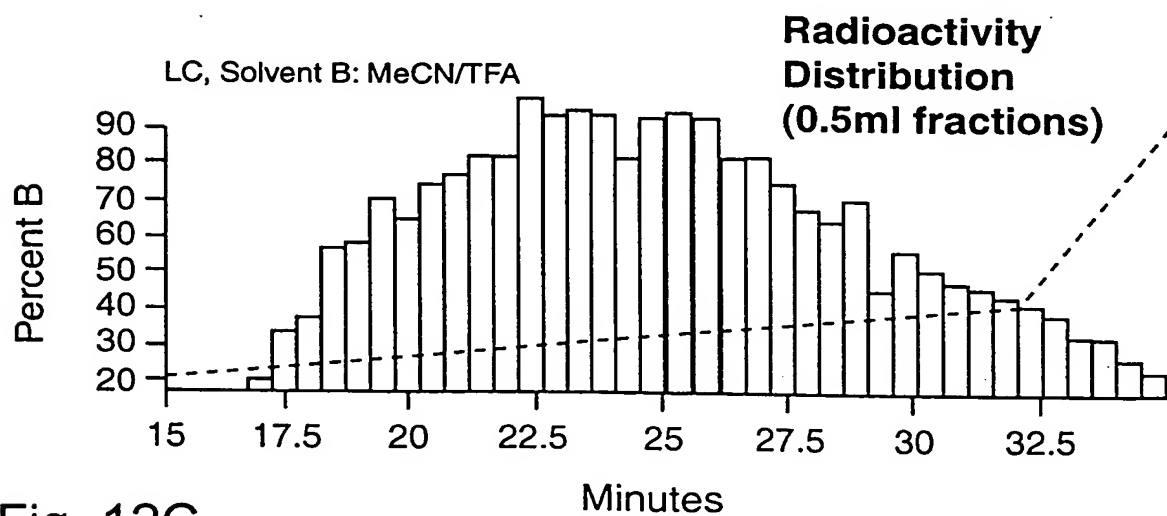


Fig. 12C

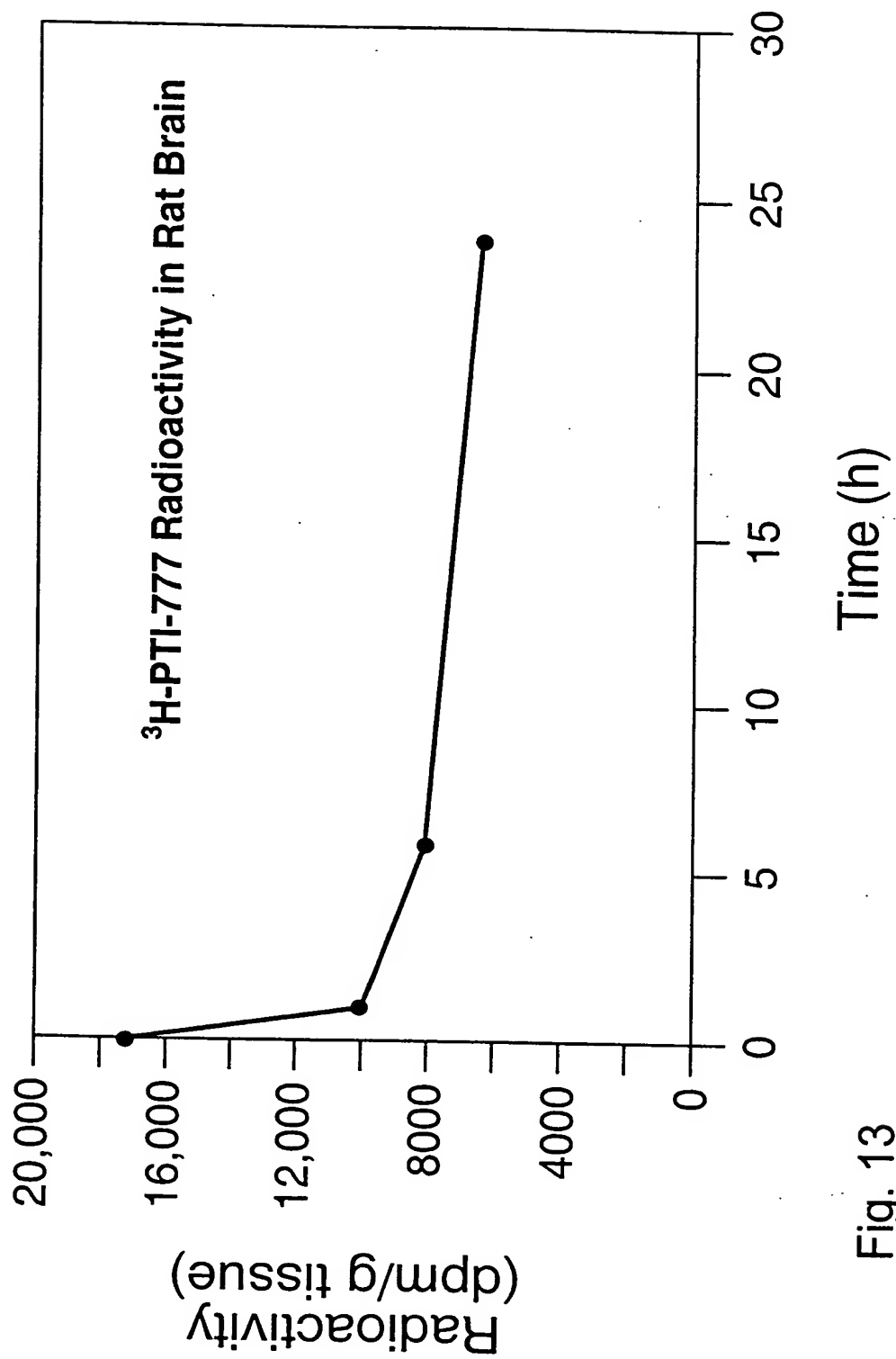


Fig. 13

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Jaguar Summed Mass Spectrum Report

File Name: G:_41.dat
Method Name:
Sample ID:
Instrument: JAGUAR

Time Run: 11/13/00 11:01:40 AM
Report Created: 11/13/00 11:03:26 AM
Operator:
Ionization Mode: ESI - positive ions

Spec # Range: 2706 - 2778

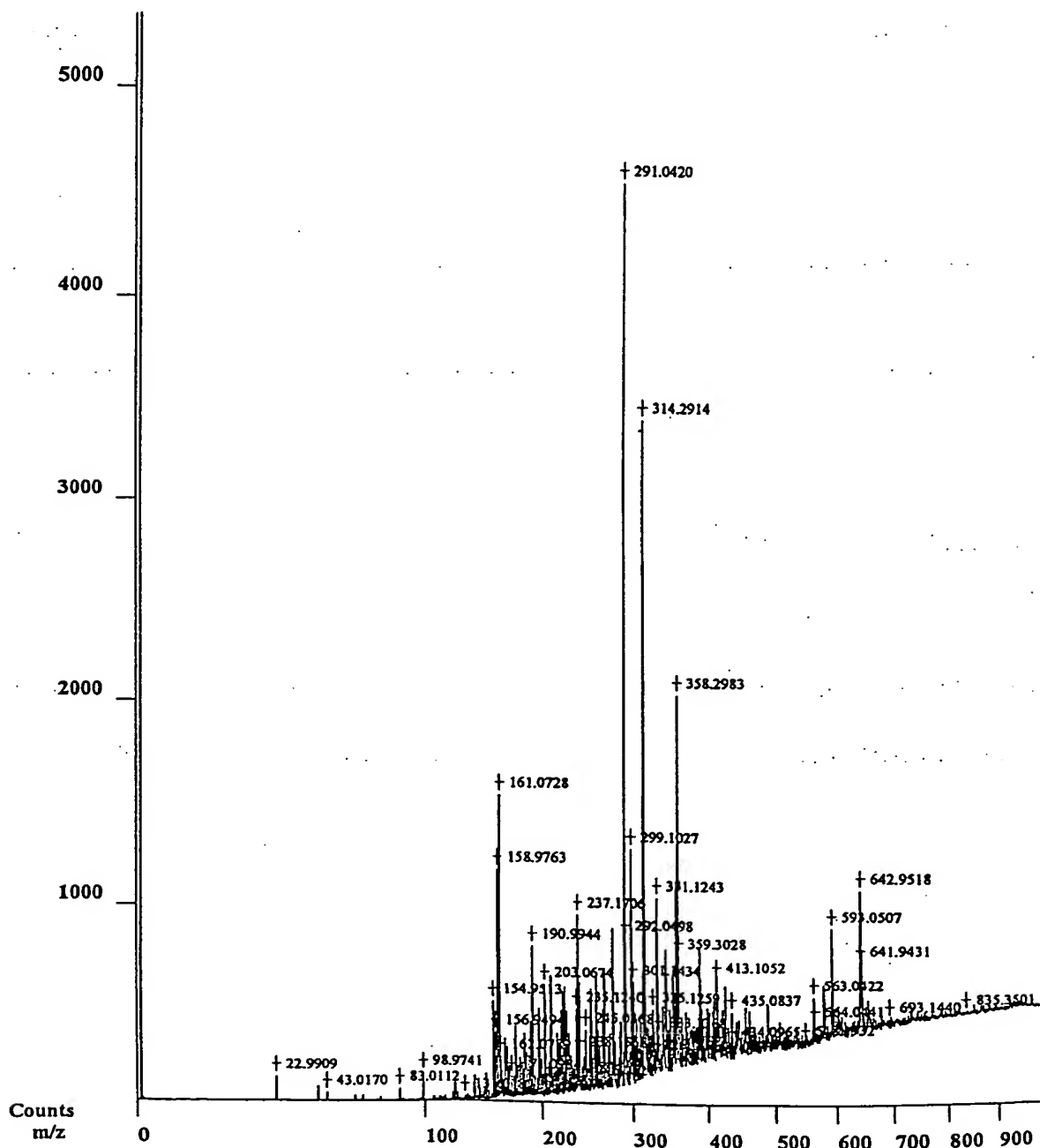


Fig. 14

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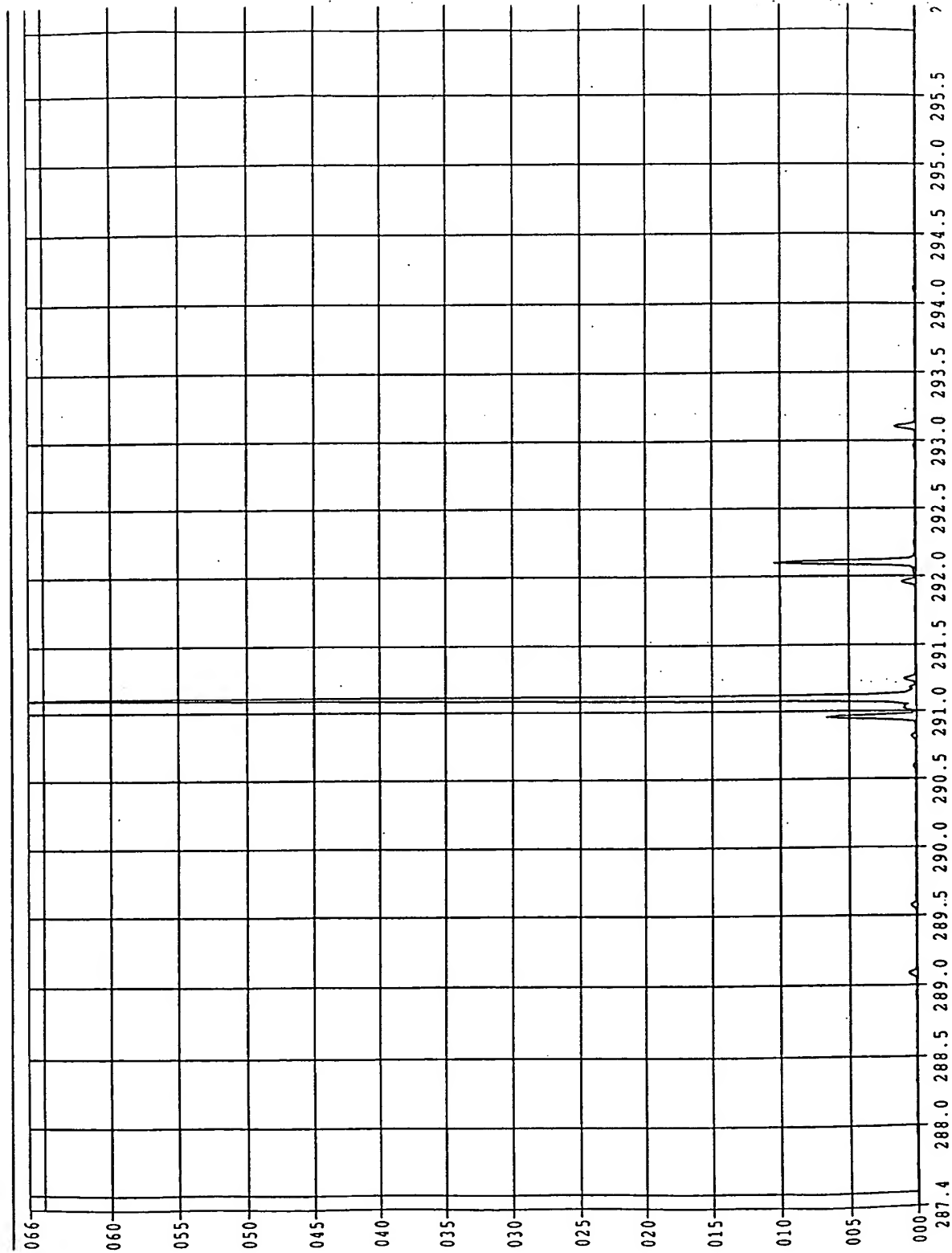


Fig. 15

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(Mass Spectrum)
Data : BYU_A776
Sample: 1
Note :
Inlet : Direct Ion Mode : EI+
Spectrum Type : Normal Ion (MF-Linear)
RT : 0.64 min Scan# : (5,8)
BP : m/z 140.0000 Int. : 135.19
Output m/z range : 50.0000 to 362.6855 Cut Level : 0.00 %

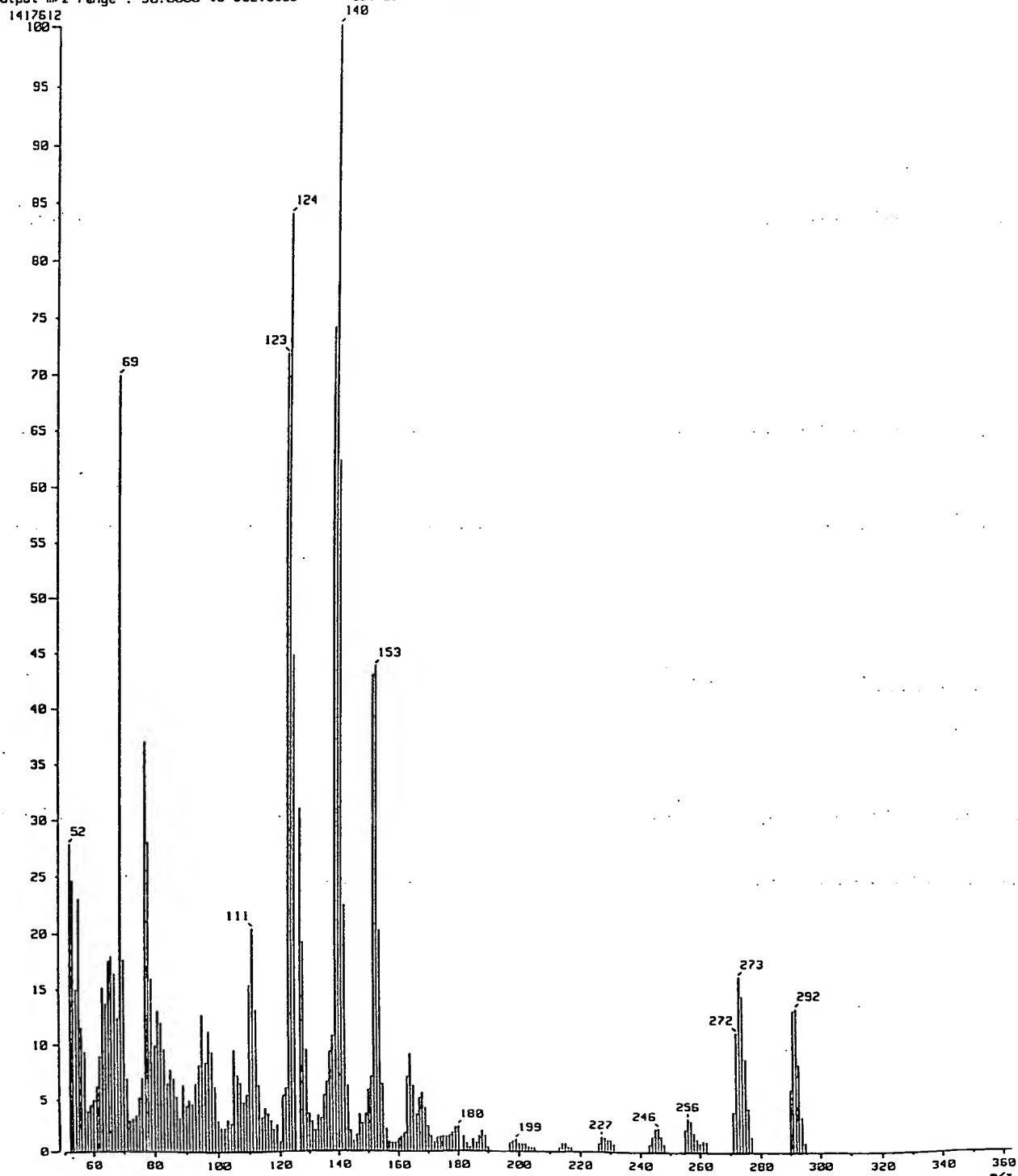


Fig. 16

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(Mass Spectrum)
Data : BYU_R349
Sample: acylated 1
Note : ThioGlycerol & Na
Inlet : Direct Ion Mode : FAB+
Spectrum Type : Normal Ion (MF-Linear)
RT : 0.08 min Scan# : (1,4)
BP : m/z 523.0000 Int. : 322.38
Output m/z range : 240.0000 to 642.3739 Cut Level : 0.00 %
3405255

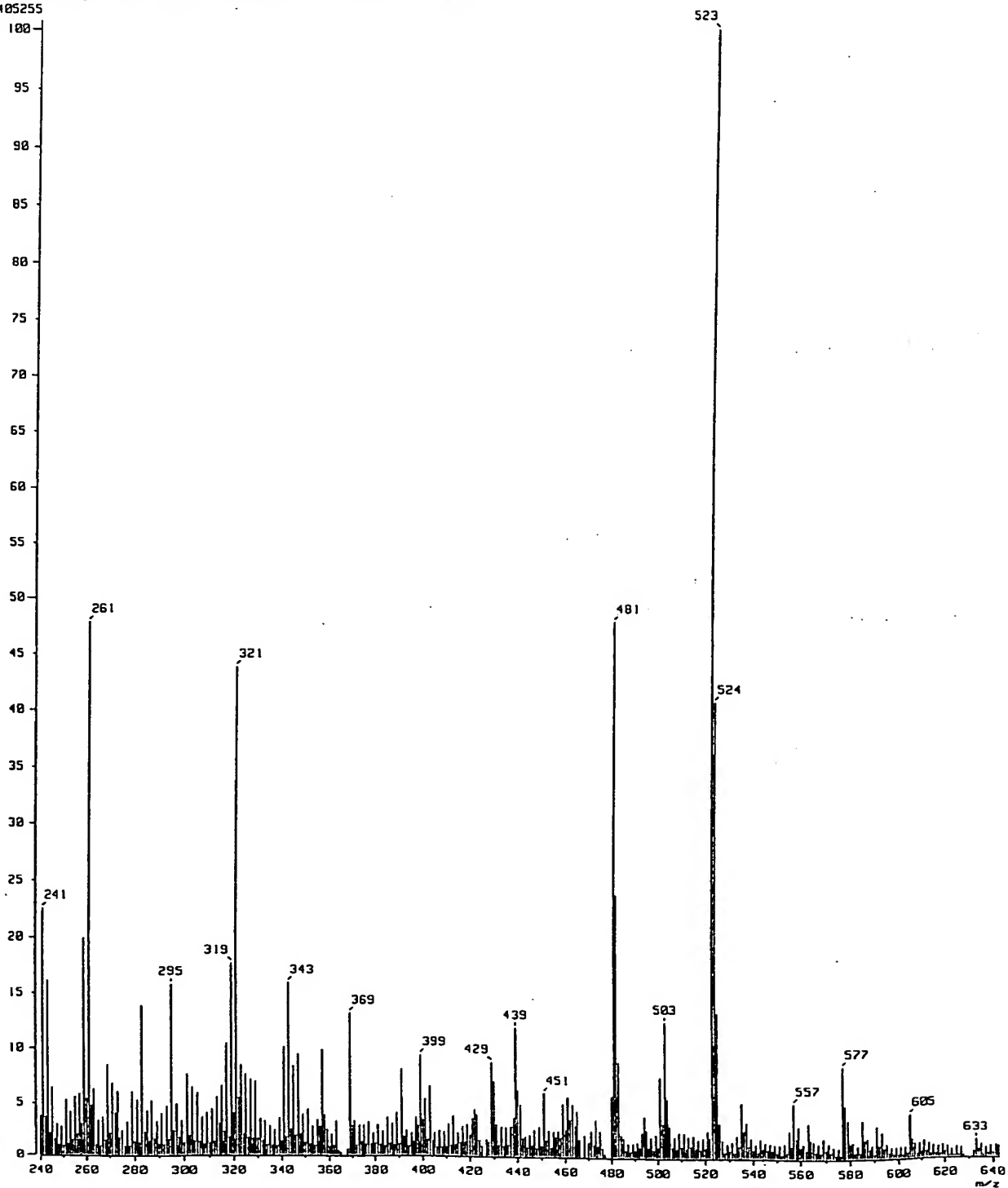


Fig. 17

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(Mass Spectrum)
Data : BYU R373
Sample: acylated-1
Note :
Inlet : Direct Ion Mode : EI+
Spectrum Type : Normal Ion (MF-Linear)
RT : 0.89 min Scan# : (7,9)-k((3,4))(k=1.0)
BP : m/z 398.0000 Int. : 68.70
Output m/z range : 50.0000 to 564.0950 Cut Level : 0.00 %

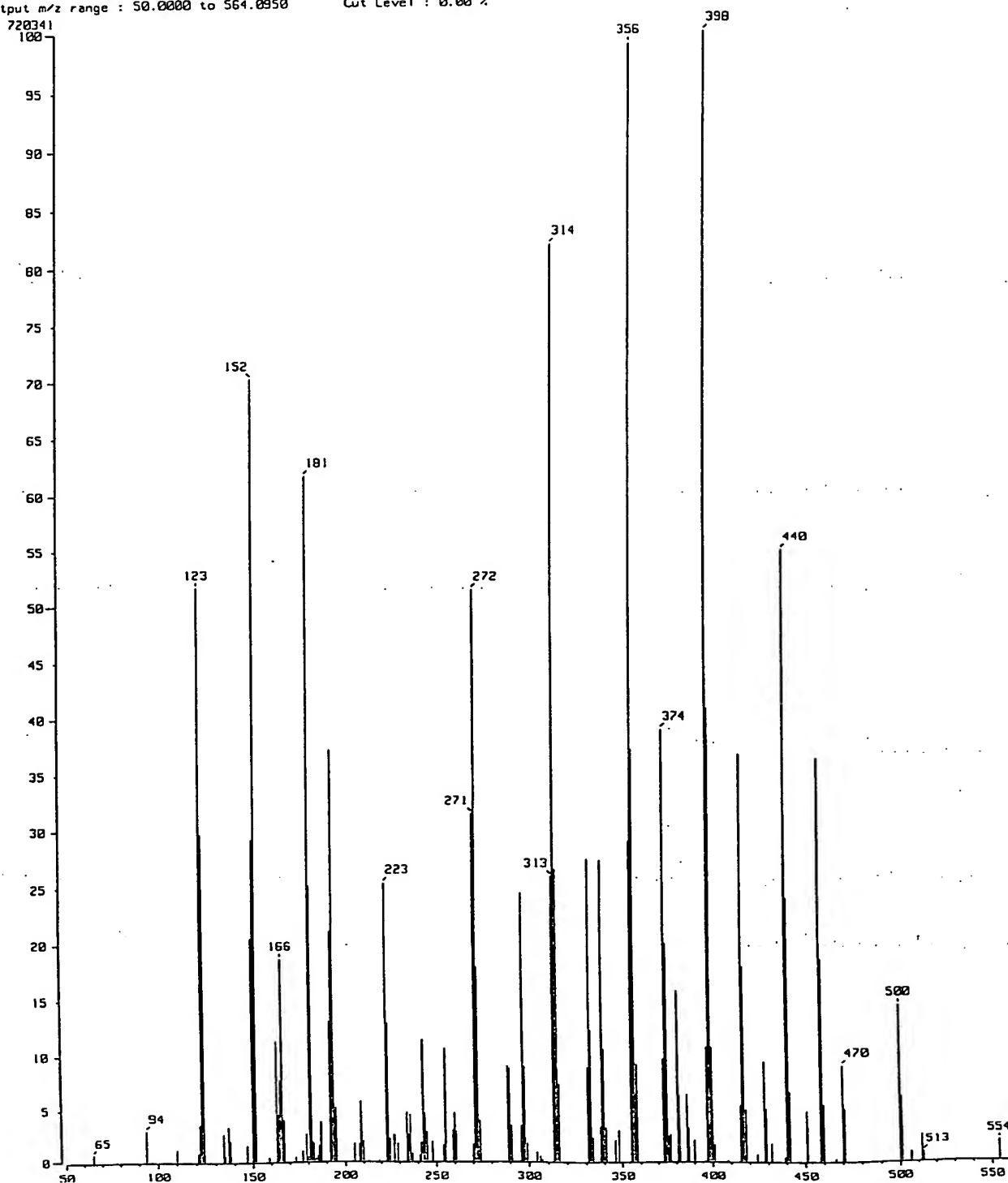


Fig. 18

H1 of sample.

Pulse Sequence: s2pul

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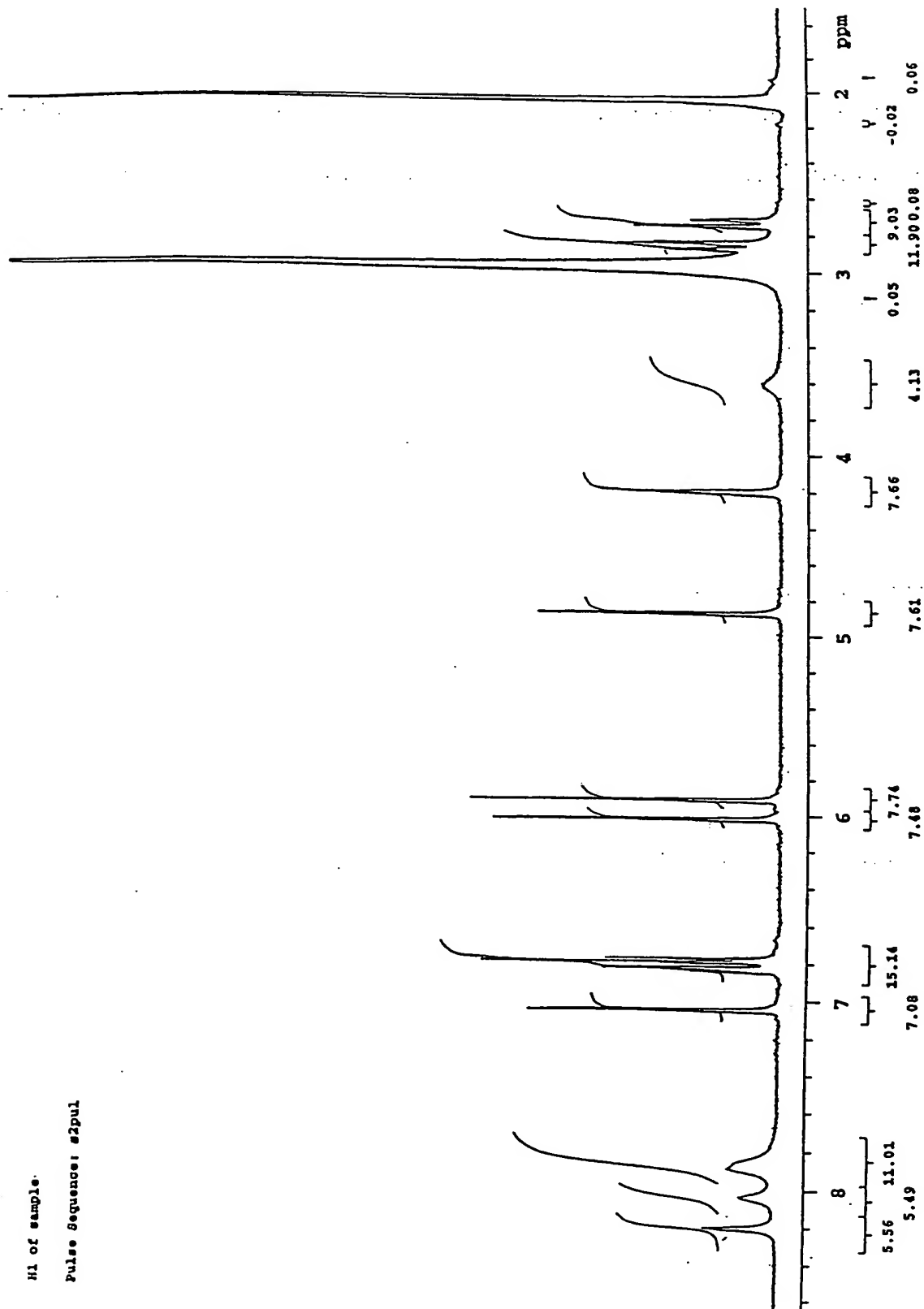


Fig. 19

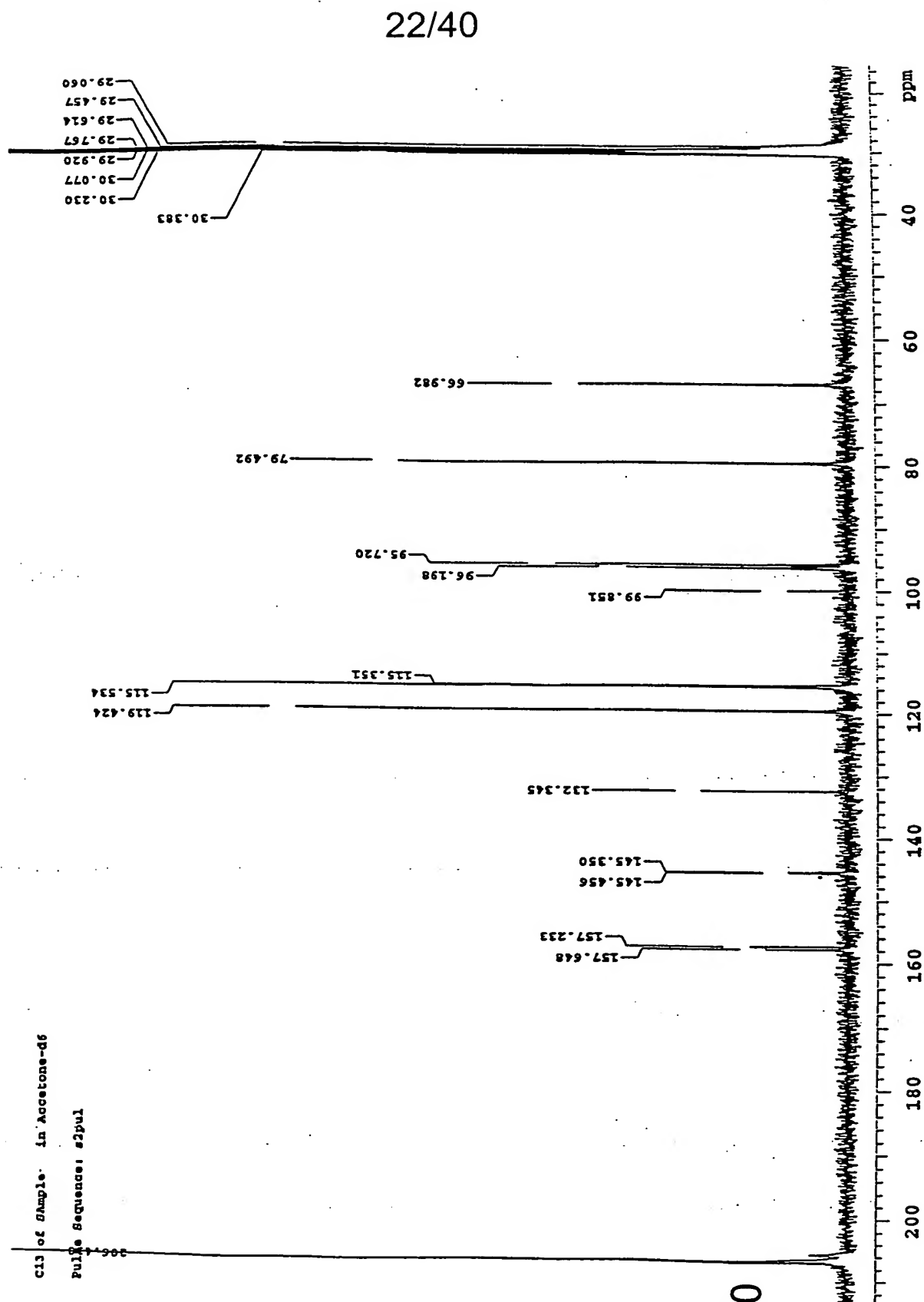


Fig. 20

H1
Pulse Sequence: s2pul

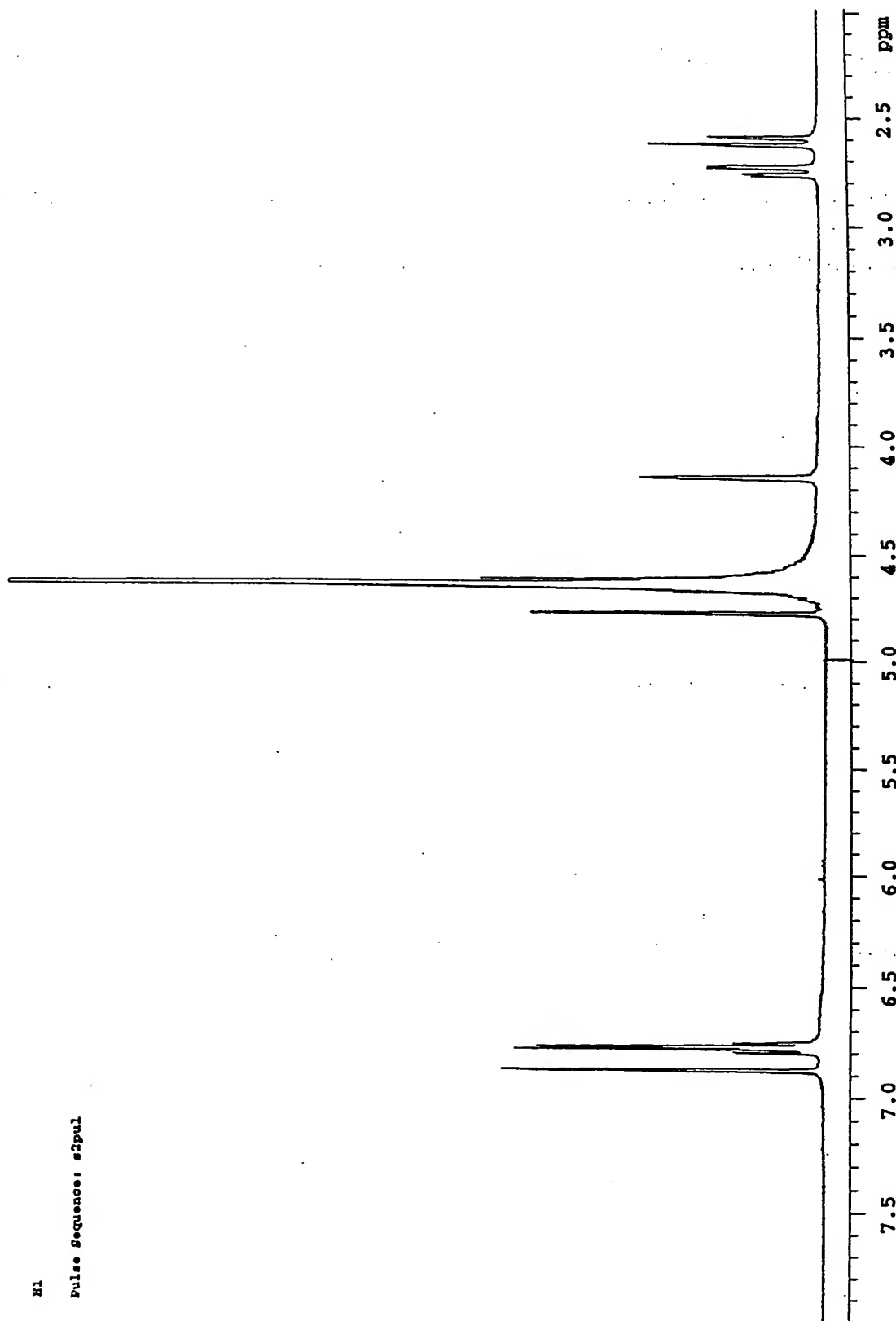


Fig. 21

C13 of Sample in D2O.

Pulse Sequence: zgpg30

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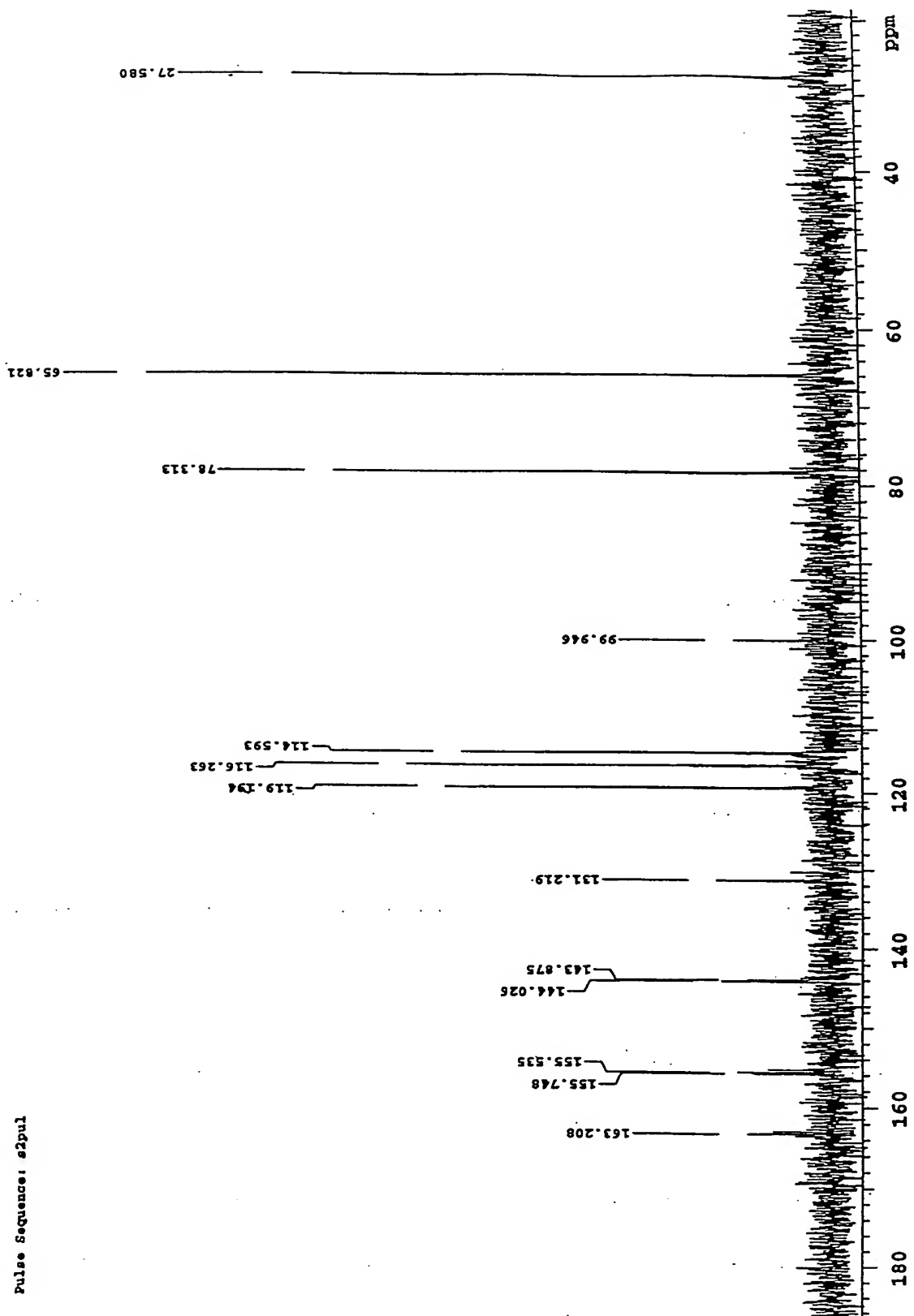
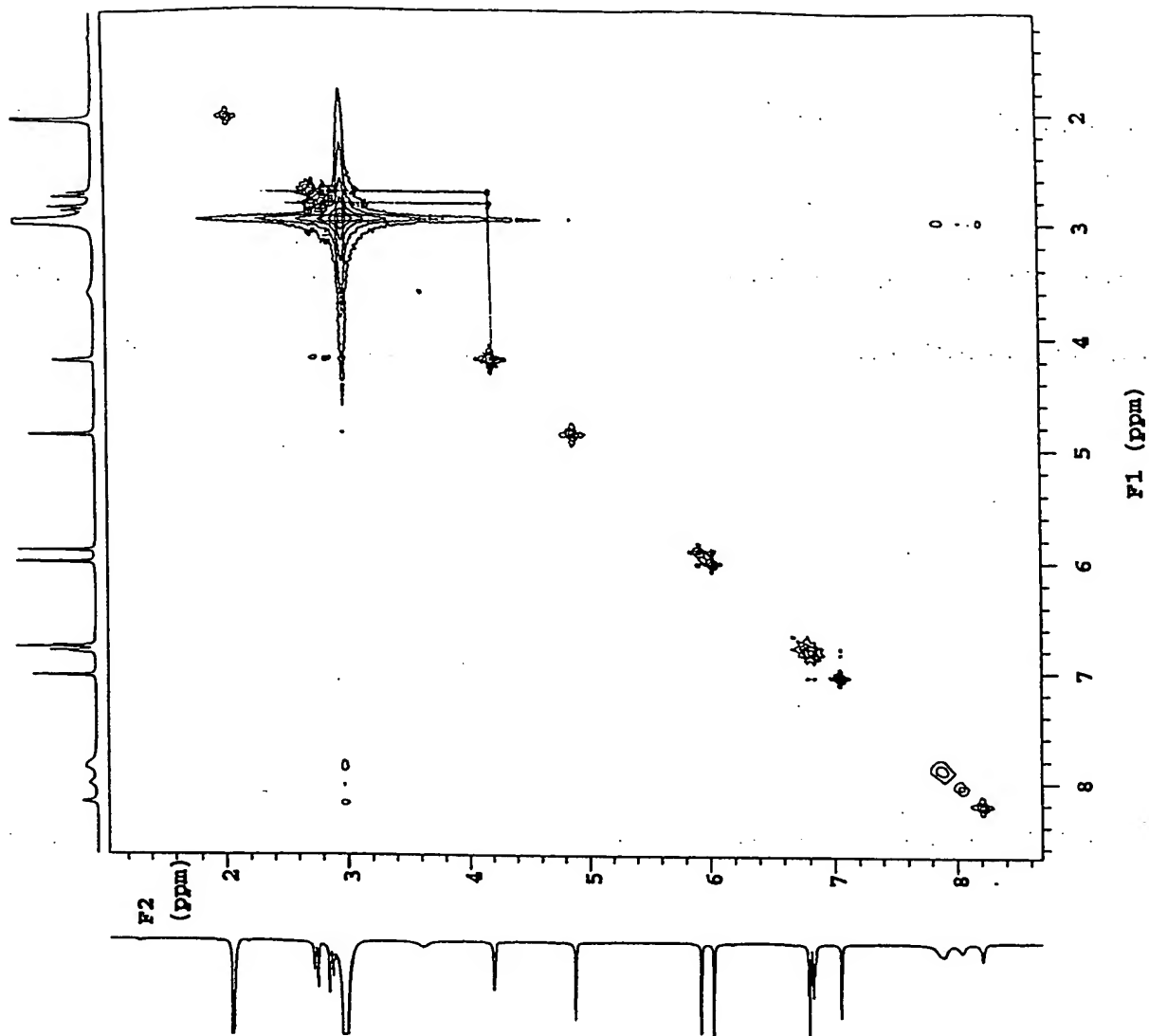


Fig. 22

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COSY of Sample
 Pulse Sequence: relayh
 Solvent: Acetone
 Ambient temperature
 UNITY-500 "nmr500"
 PULSE SEQUENCE: relayh
 COSY 90-90
 Acq. time 0.302 sec
 Width 3817.5 Hz
 2D Width 3817.5 Hz
 16 repetitions
 159 increments
 OBSERVE H1, 499.881428 MHz
 DATA PROCESSING
 F1 DATA PROCESSING
 Line broadening 0.3 Hz
 FT size 1024 x 1024
 Total time 14 min, 12 sec

Fig. 23

COSY

Pulse Sequence: relayh

Solvent: D2O

Ambient temperature

UNITY-500 "nmr500"

PULSE SEQUENCE: relayh

Relax. delay 0.500 sec

COSY 90-90

Acq. time 0.178 sec

Width 2882.3 Hz

2D Width 2882.3 Hz

8 repetitions

120 increments

OBSERVE H1, 499.8801324 MHz

DATA PROCESSING

Line broadening 0.1 Hz

F1 DATA PROCESSING

Line broadening 0.3 Hz

F1 size 1024 x 1024

Total time 11 min, 23 sec

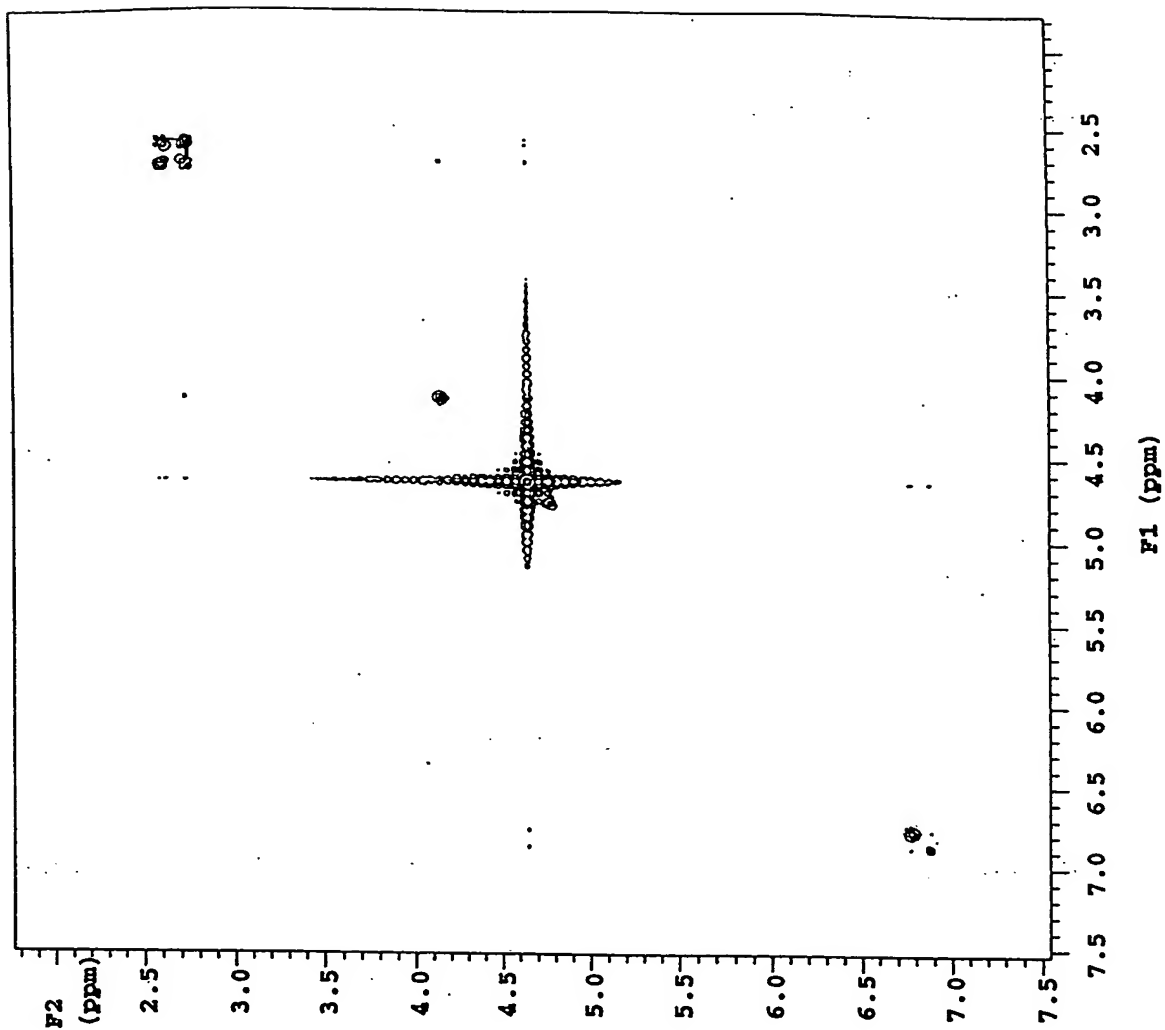
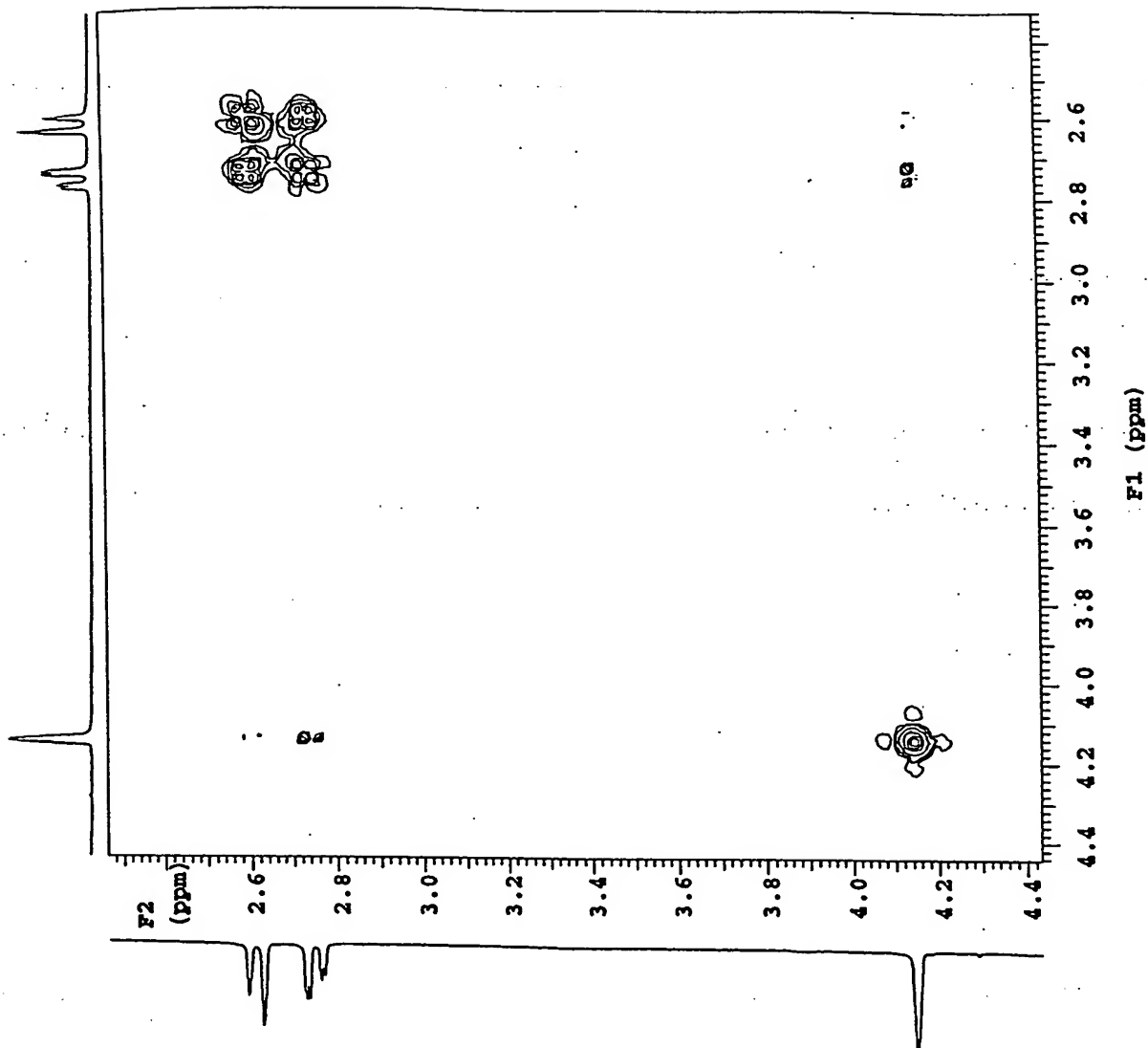


Fig. 24A

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COSY

Pulse Sequence: relayh

Solvent: D2O

Ambient temperature

UNITY-500 "nmr500"

PULSE SEQUENCE: relayh

Relax. delay 0.500 sec

COSY 90-90

Acq. time 0.178 sec

Width 2882.3 Hz

2D Width 2882.3 Hz

8 repetitions

120 increments

OBSERVE H1, 499.8801324 MHz

DATA PROCESSING

Line broadening 0.1 Hz

F1 DATA PROCESSING

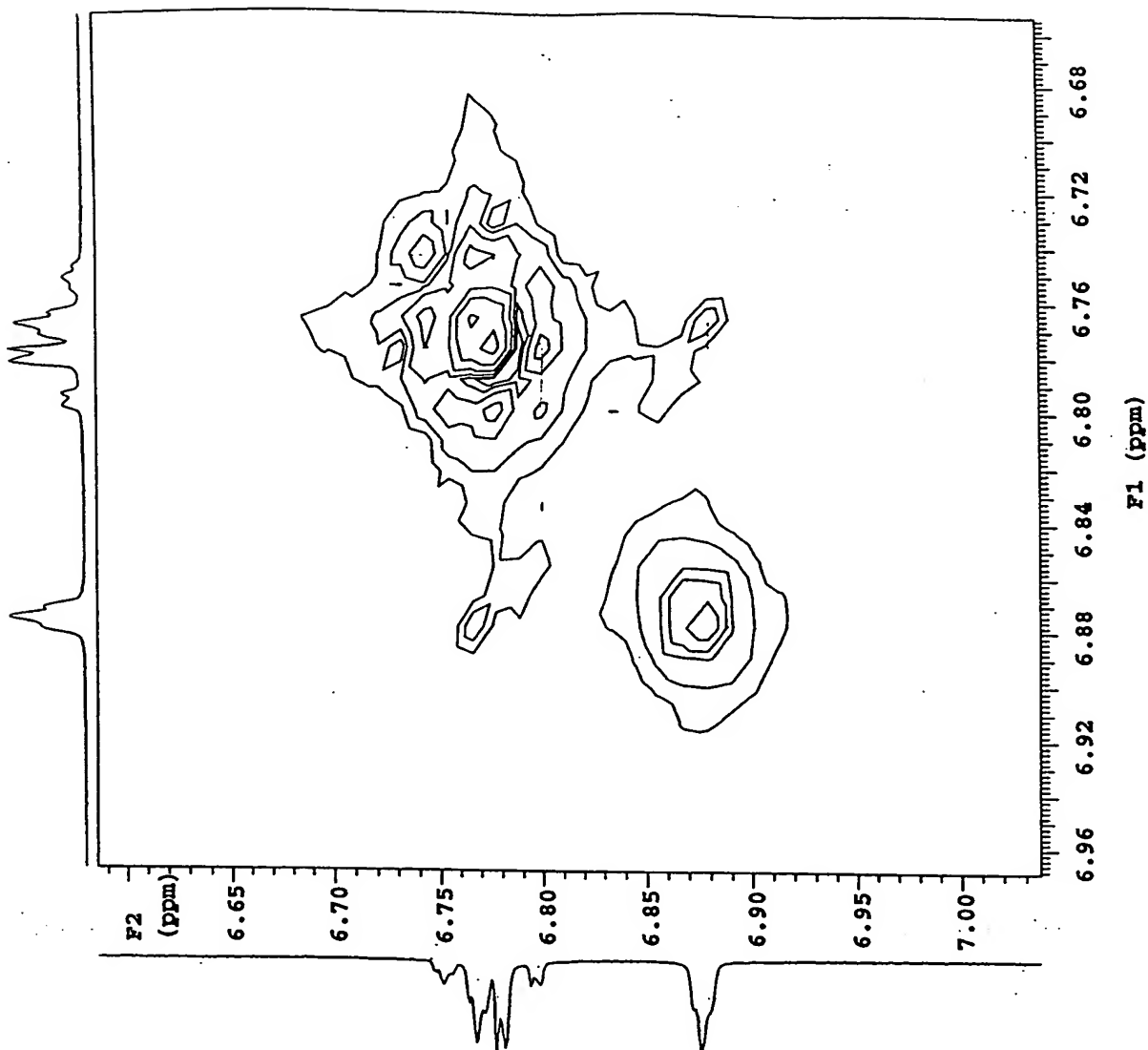
Line broadening 0.3 Hz

FT size 1024 x 1024

Total time 11 min, 23 sec

Fig. 24B

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COSY

Pulse Sequence: relayh

Solvent: D2O

Ambient temperature

UNITY-500 "nmr500"

PULSE SEQUENCE: relayh

Relax. delay 0.500 sec

COSY 90-90

Acq. time 0.178 sec

Width 2882.3 Hz

2D Width 2882.3 Hz

8 repetitions

120 increments

OBSERVE H1, 499.8801324 MHz

DATA PROCESSING

Line broadening 0.1 Hz

F1 DATA PROCESSING

Line broadening 0.3 Hz

FT size 1024 x 1024

Total time 11 min, 23 sec

Fig. 24C

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H1 of 1 Acetylated in CDCl3

Pulse Sequence: s2pul

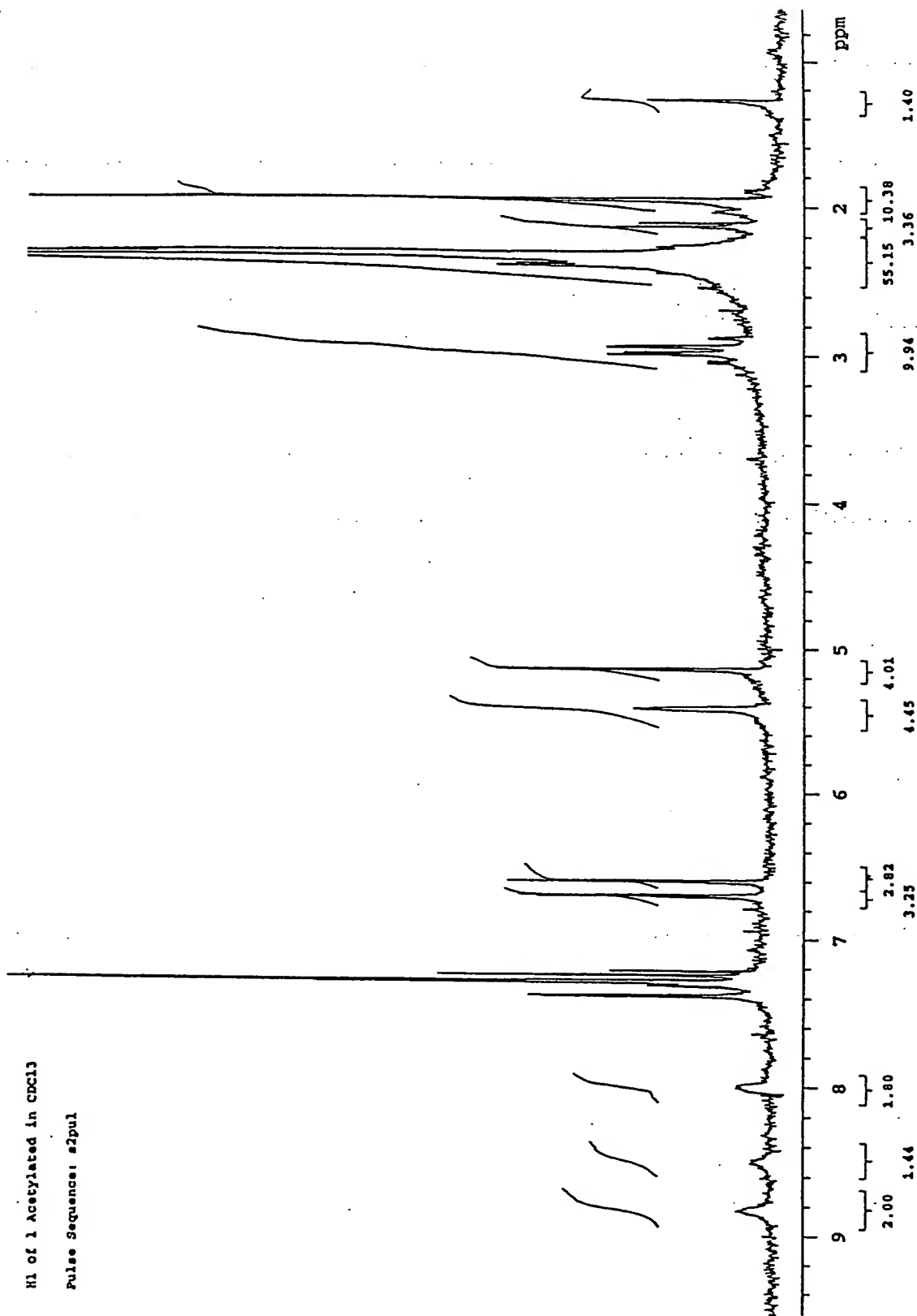
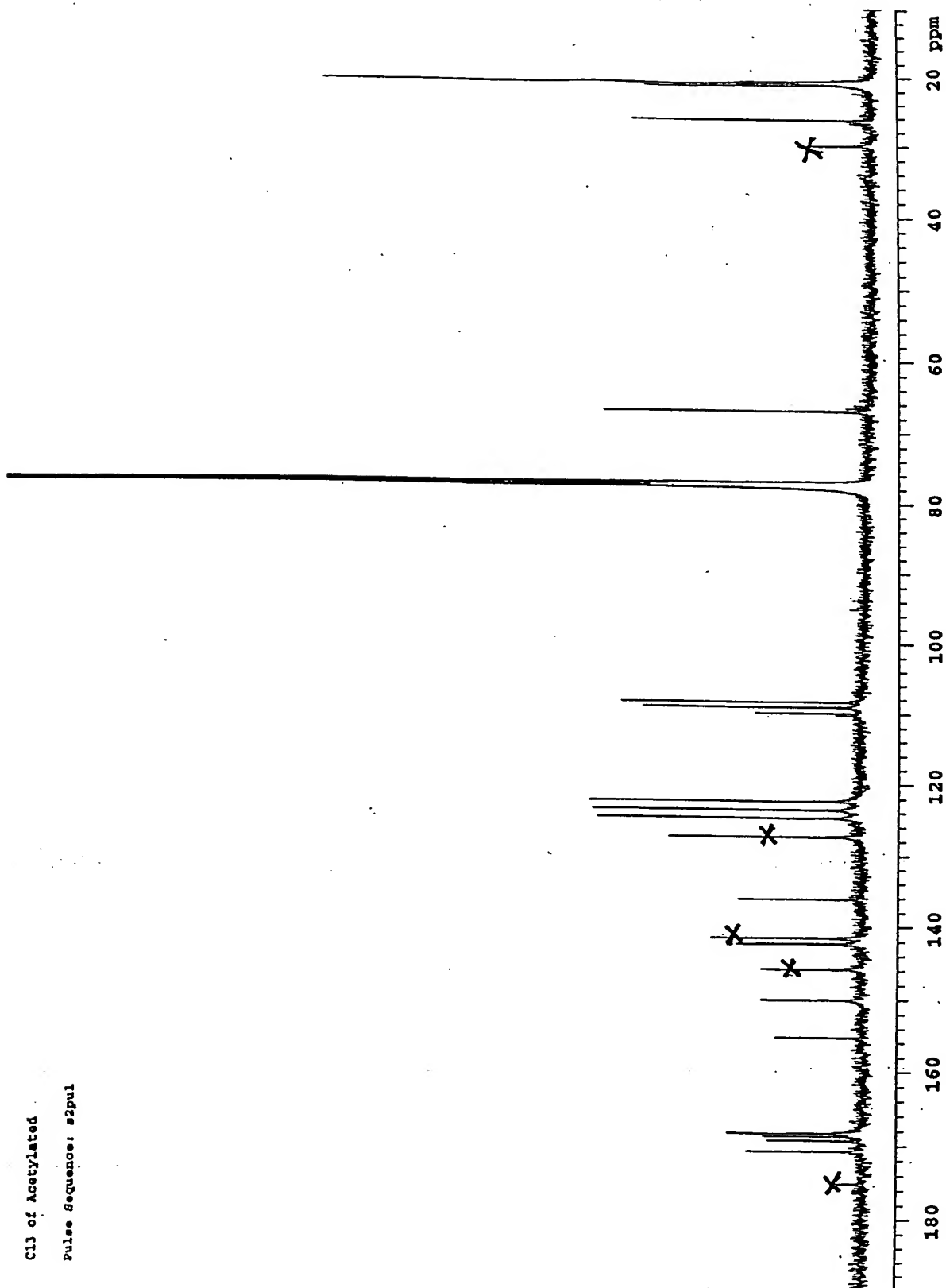


Fig. 25

C13 of Acetylated
Pulse Sequence: s2pul

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X=pyridinium acetate

Fig. 26

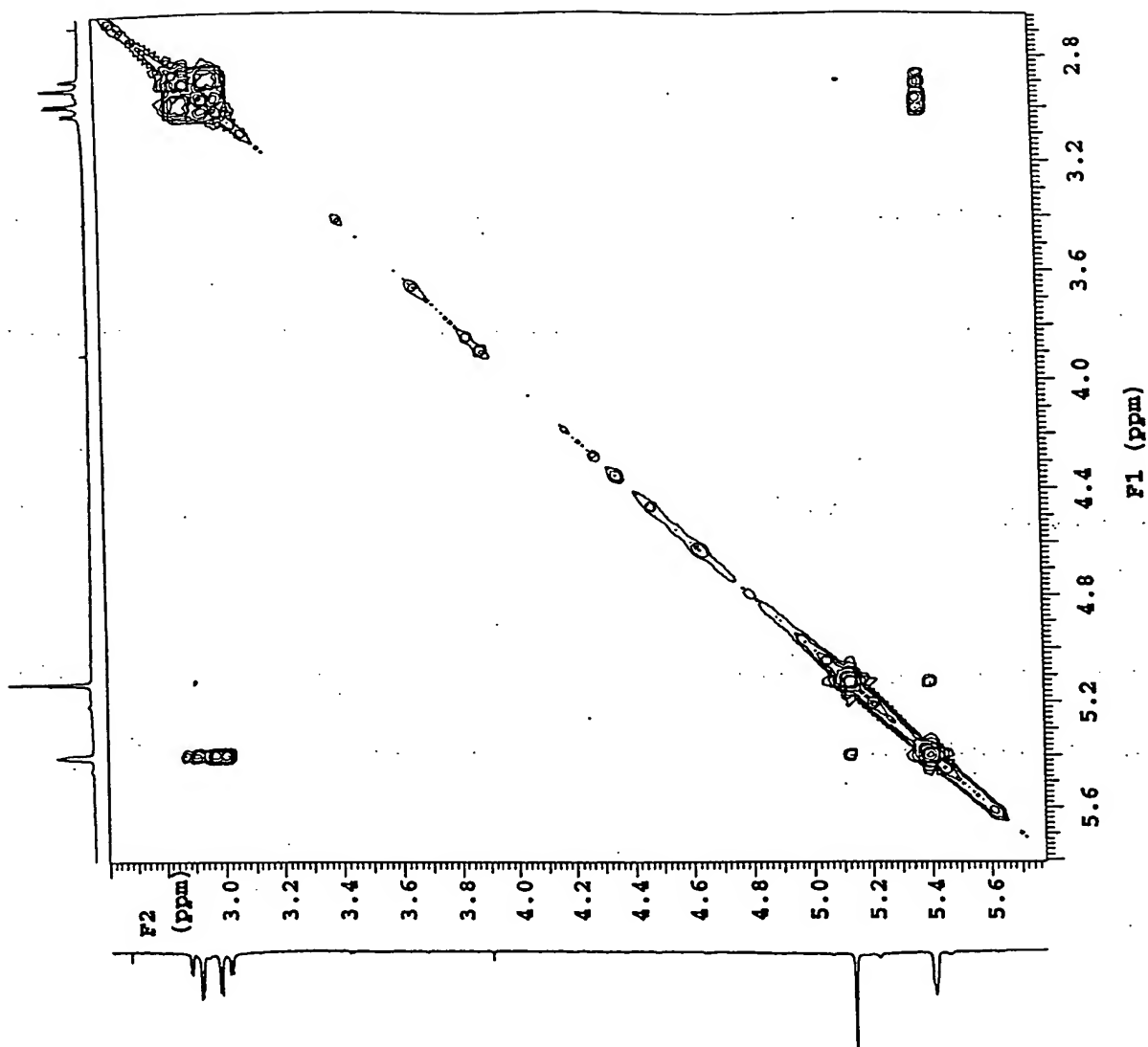


Fig. 27

HETCOR of Acetylated . in CDCl₃.

Pulse Sequence: hetcor

Solvent: CDCl₃

Temp. 26.0 C / 299.1 K

User: 1-14-87

INOVA-500 ¹Hmr500

Relax. delay 1.000 sec

Acq. time 0.082 sec

Width 25000.0 Hz

2D Width 4614.9 Hz

208 repetitions

256 increments

OBSERVE C13, 125.7033376 MHz

DECOUPLE H1, 499.9160715 MHz

Power 36 dB

on during acquisition

off during delay

WALTZ-16 modulated

DATA PROCESSING

Line broadening 2.0 Hz

F1 DATA PROCESSING

Line broadening 0.3 Hz

FT size 4096 x 1024

Total time 16 hr, 46 min, 21 sec

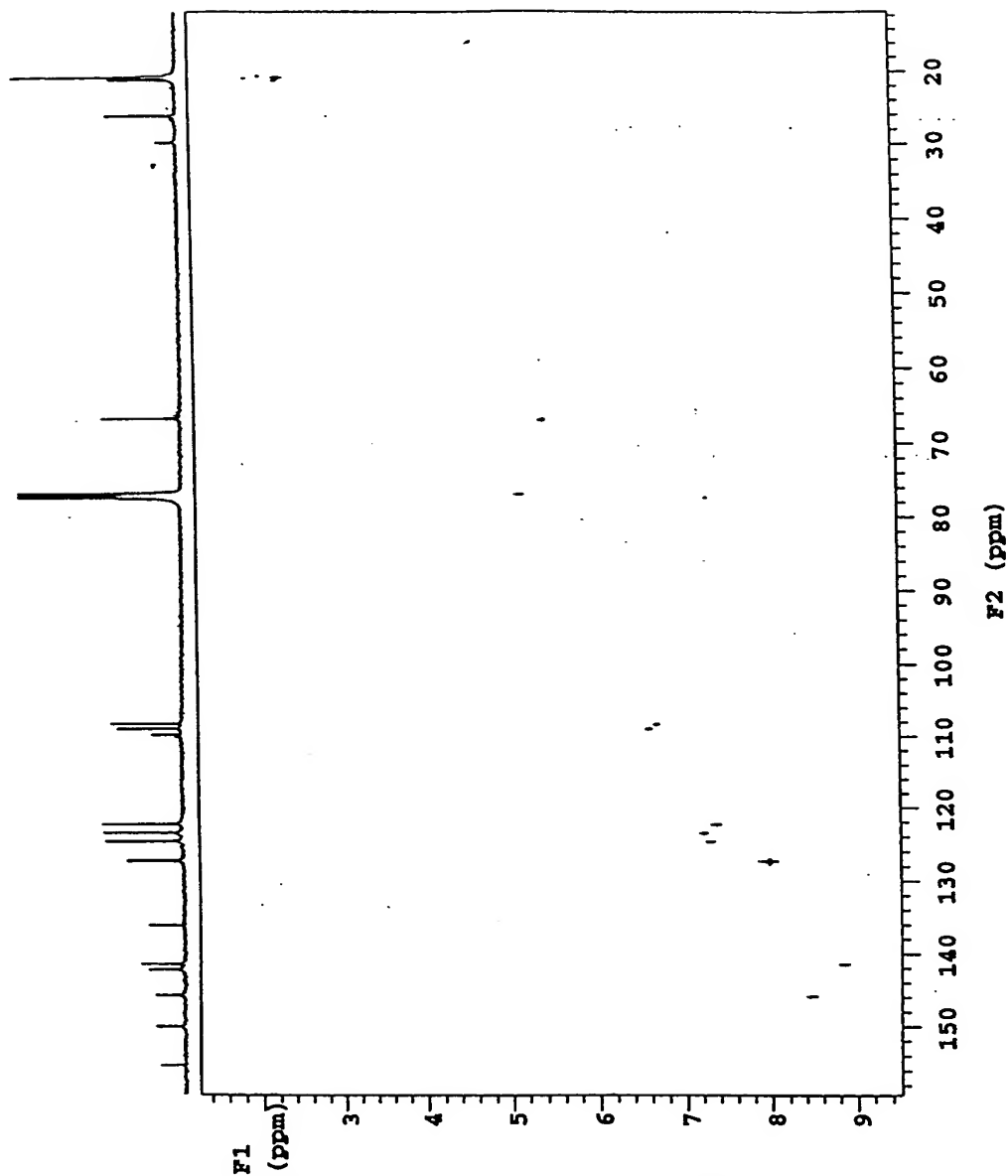


Fig. 28A

HETCOR of Acetylated in CDCl₃.

X=pyridinium acetate

Pulse Sequence: hetcor

Solvent: CDCl₃

Temp. 26.0 C / 299.1 K

User: 1-14-87

INOVA-500 "nmr500"

Relax. delay 1.000 sec

Acq. time 0.082 sec

Width 25000.0 Hz

2D Width 4614.9 Hz

208 repetitions

256 increments

OBSERVE C13, 125.7033376 MHz

DECOUPLE H1, 499.9160715 MHz

Power 38 dB

on during acquisition

off during delay

WALTZ-16 modulated

DATA PROCESSING

Line broadening 2.0 Hz

F1 DATA PROCESSING

Line broadening 0.3 Hz

FT size 4096 x 1024

Total time 16 hr, 46 min, 21 sec

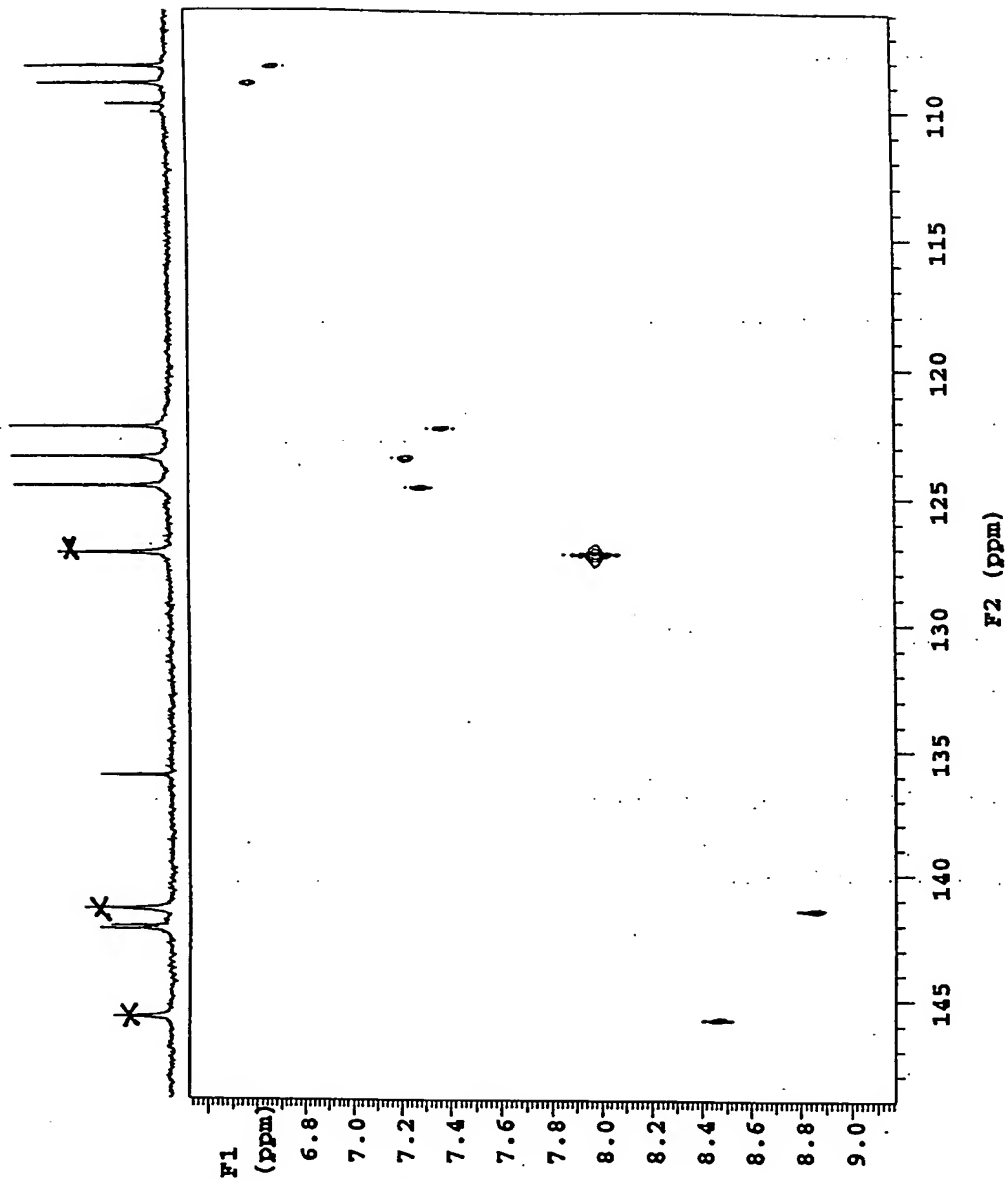


Fig. 28B

HETCOR of Acetylated . in CDCl₃.

Pulse Sequence: hetcor

Solvent: CDCl₃

Temp. 26.0 C / 299.1 K

User: 1-14-87

INNOVA-500 "nmr500"

Relax. delay 1.000 sec

Acq. time 0.082 sec

Width 25000.0 Hz

2D Width 4614.9 Hz

208 repetitions

256 increments

OBSERVE C13, 125.7033376 MHz

DECOUPLE H1, 499.9160715 MHz

Power 38 dB

on during acquisition

off during delay

WALTZ-16 modulated

DATA PROCESSING

line broadening 2.0 Hz

F1 DATA PROCESSING

line broadening 0.3 Hz

FT size 4096 x 1024

Total time 16 hr, 46 min, 21 sec

X=pyridinium acetate

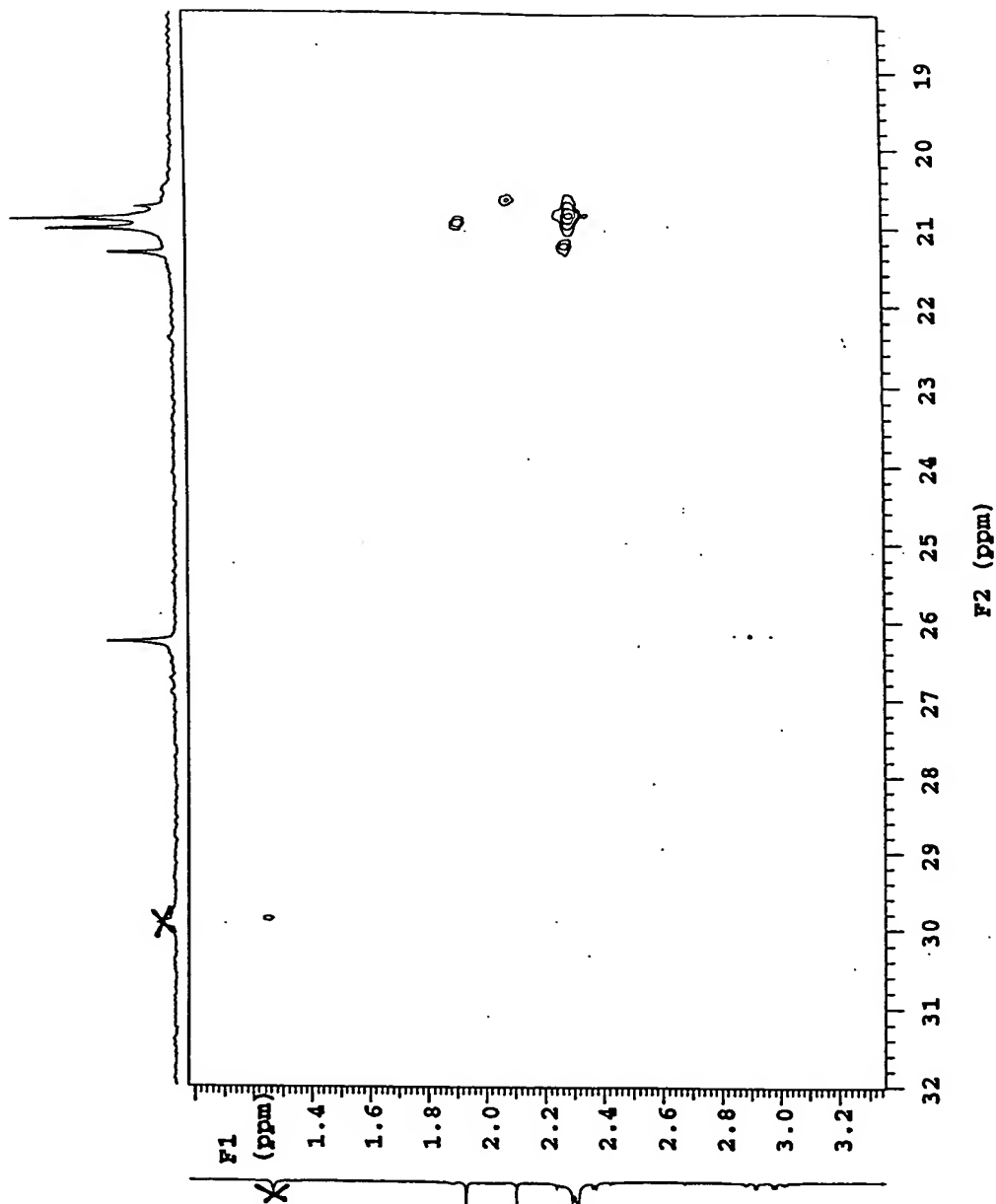


Fig. 28C

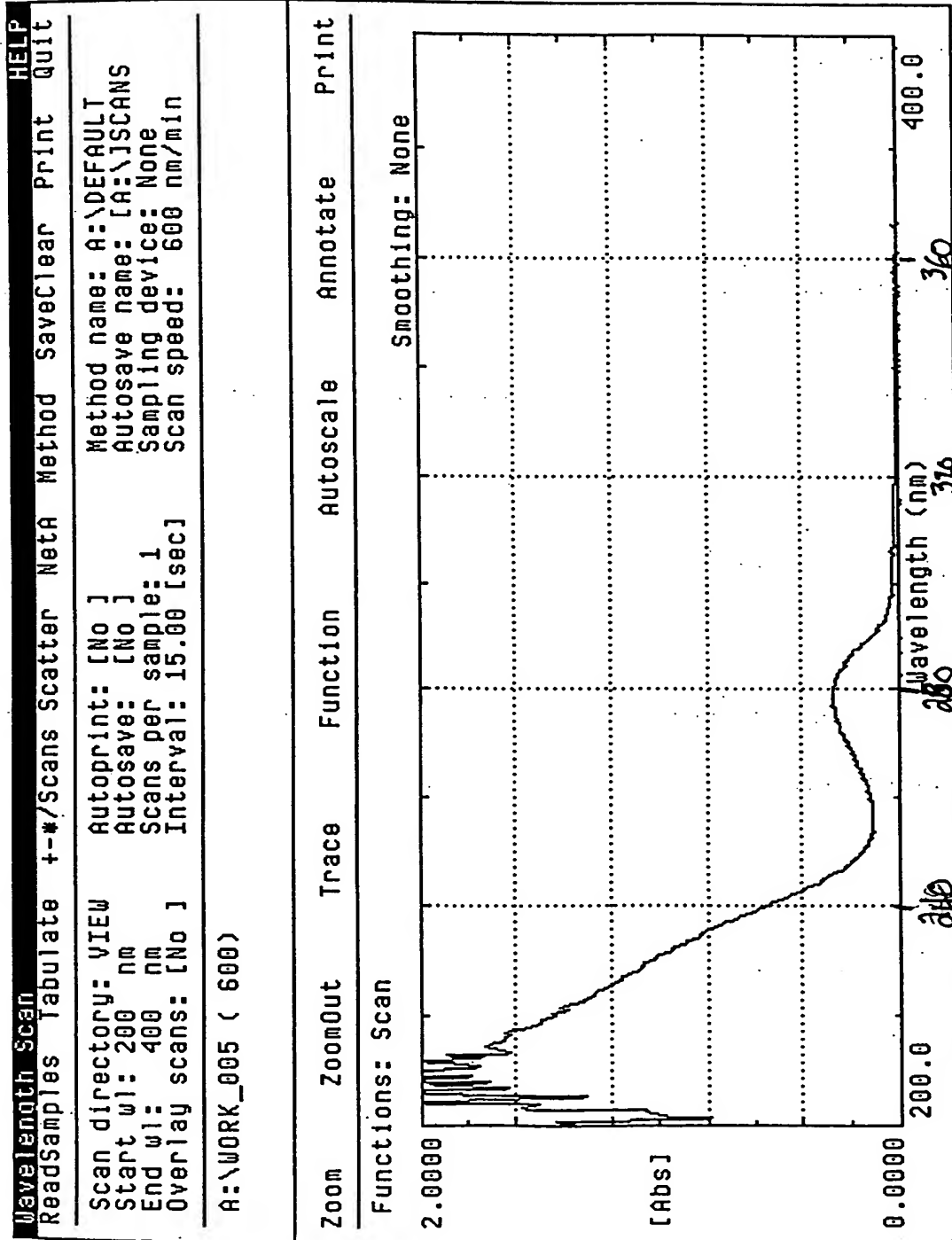


Fig. 29

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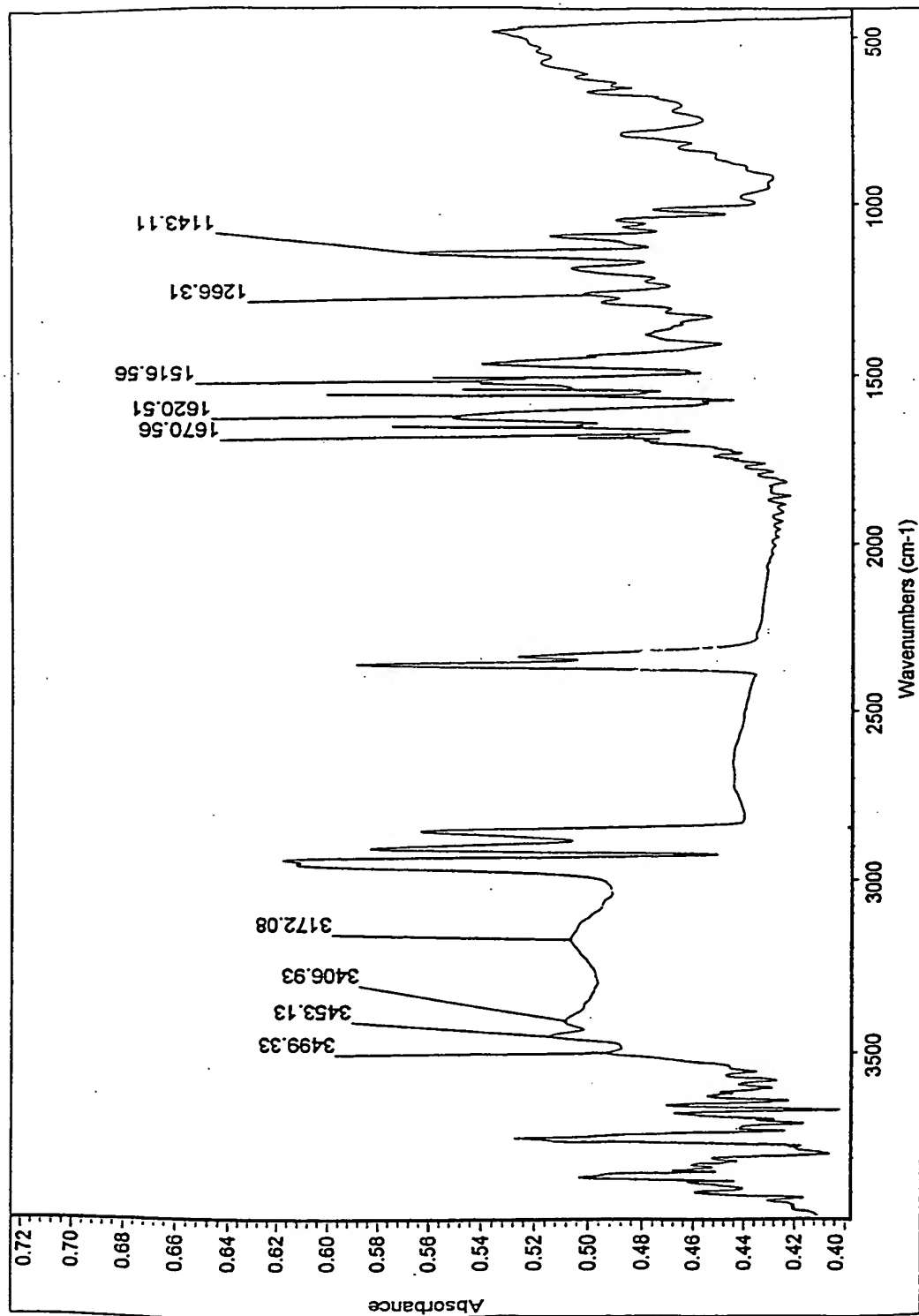


Fig. 30

**Subtraction Result: proteotech, sampleL

Scans: 32

Resolution: 4.000

Aromatic Alcohols and Phenols

Aldrich 85,523-5 CAS [490-46-0]
 (-)-Epicatechin, 97%

$C_{15}H_{14}O_6$ FT-IR: 1, 1120D
 FW 280.27
 mp 240°C d.

166.31 130.42 94.89*
 166.01 117.76* 93.81*
 155.57 114.69* 77.89*
 144.28 114.58* 64.75*
 144.23 98.31 28.07

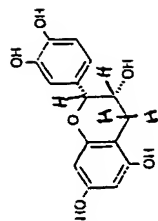
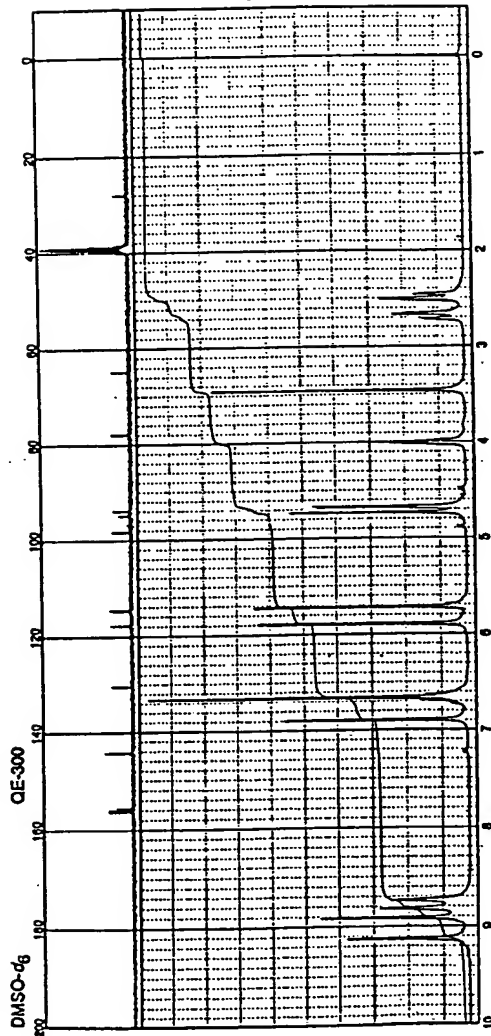


Fig. 31A

Aldrich 86,181-2
 (±)-Catechin hydrate, 98%

$C_{15}H_{14}O_6$
 FW 280.28
 mp 210°C d.

156.25 114.35*
 155.97 98.90
 155.15 94.97*
 144.66 93.70*
 130.43 80.83*
 118.24* 66.17*
 114.92* 27.70

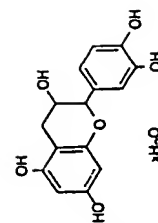
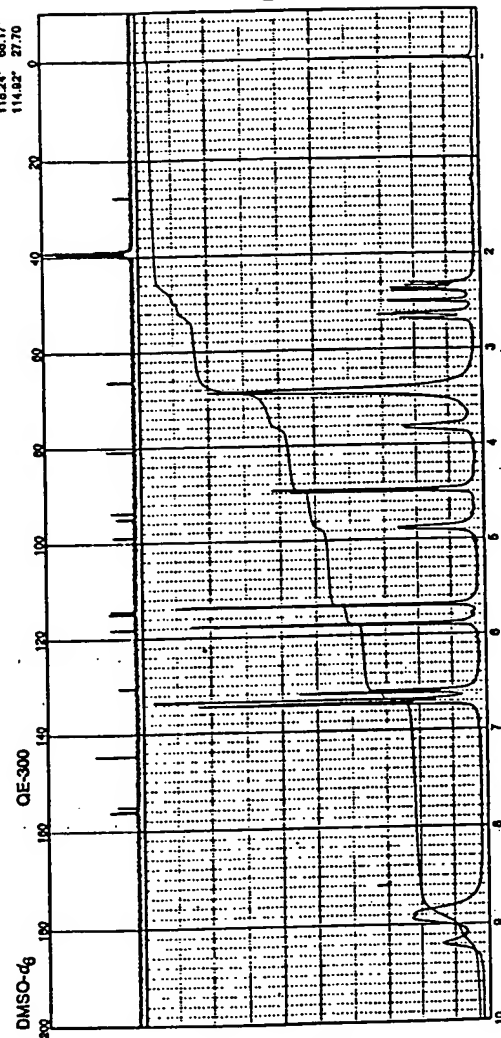


Fig. 31B

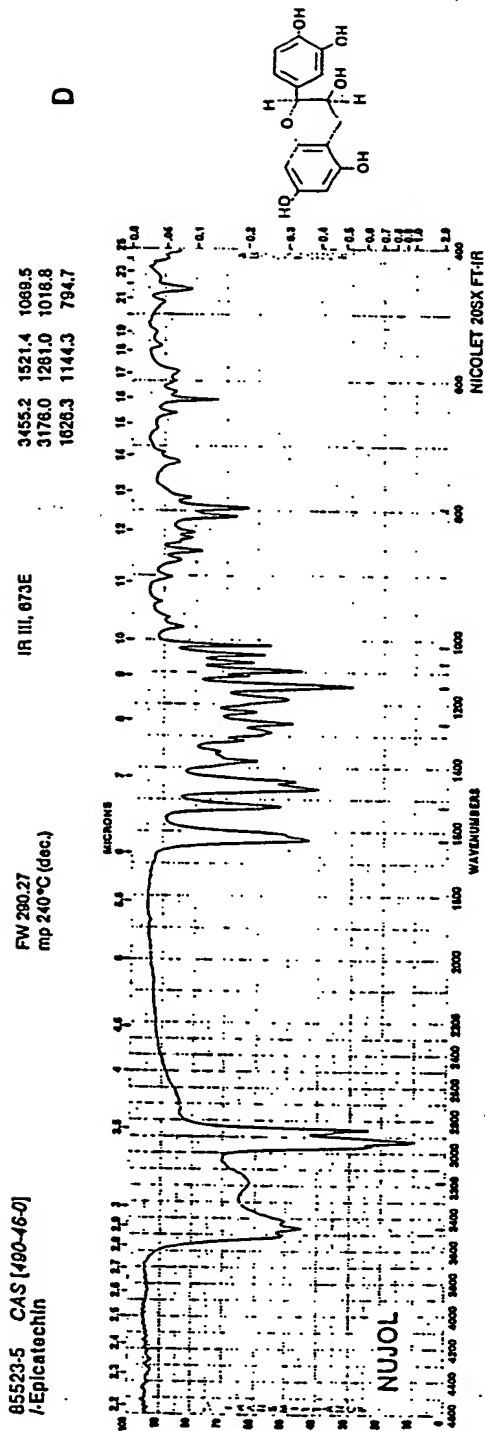


Fig. 32A

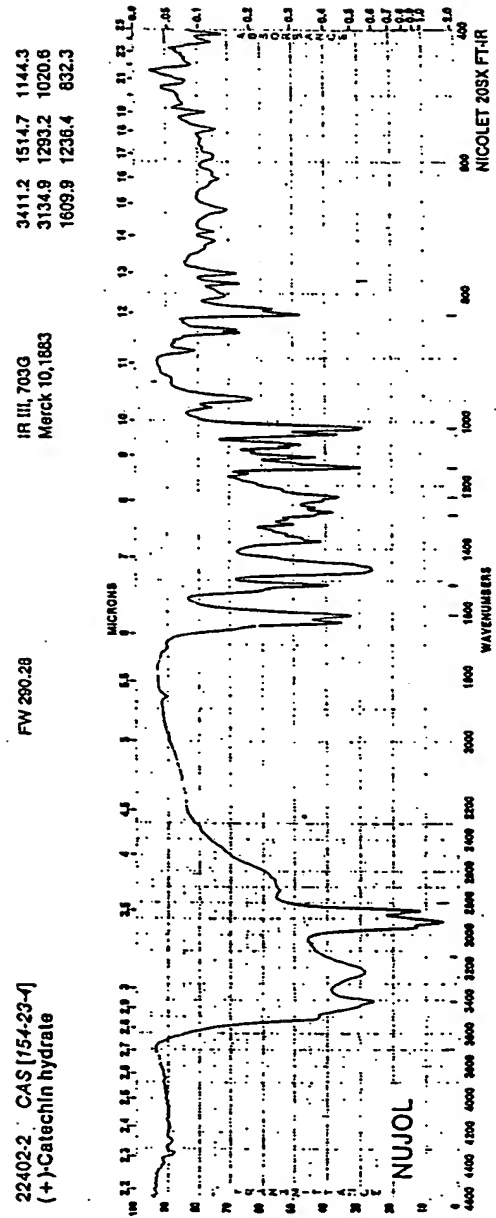


Fig. 32B

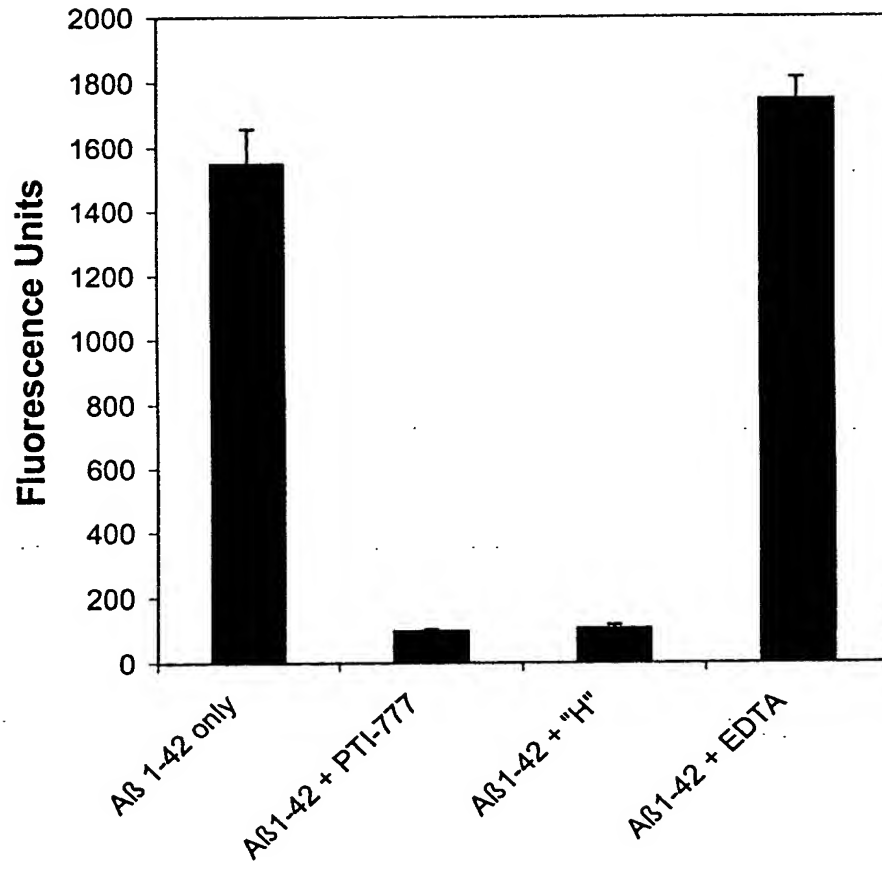


Fig. 33

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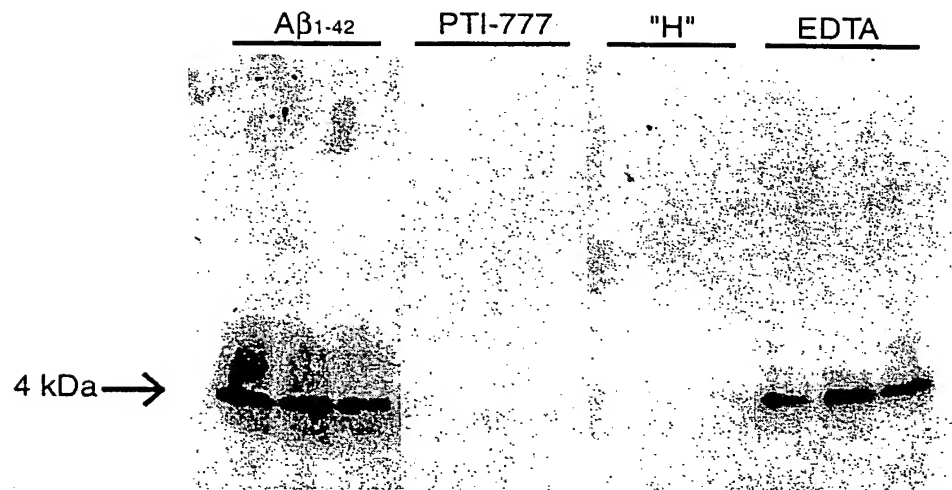


Fig. 34

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